

1 **Title:** Ascorbic acid can promote the generation and expansion of
2 neuroepithelial-like stem cells derived from hiPS/ES cells under chemically
3 defined conditions through promoting collagen synthesis

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37 **Abstract**

38 **Introduction:** Spinal cord injury (SCI) is a neurological, medically incurable
39 disorder. Human pluripotent stem cells (hPSCs) have the potential to
40 generate neural stem/progenitor cells (NS/PCs) which hold promise in
41 therapy for SCI by transplantation. In our study, we aimed to establish a
42 chemically defined culture system by using serum-free medium and
43 ascorbic acid (AA) to generation and expansion of long-term self-renewing
44 neuroepithelial-like stem cells (It-NES cells) differentiated from hPSCs
45 effectively and stably.

46 **Methods:** We induce hESC/iPSC to neurospheres by using a newly
47 established induction system in vitro in our study. And It-NES cells derived
48 from hESCs/iPSCs-neurospheres using two induction systems, including
49 conventional N2 medium with gelatin-coated (coated) and N2+AA medium
50 without pre-coated (AA) were characterized by reverse
51 transcription-polymerase chain reaction (RT-PCR) analysis and
52 immunocytochemistry staining. Subsequently, It-NES cells were induced to
53 neurons and the microelectrode array (MEA) recording system was used to
54 evaluate the functionality of neurons differentiated from It-NES cells.

55 Moreover, the mechanism of AA-induced It-NES cells was explored through
56 RNA-seq and the use of inhibitors.

57 **Results:** HESCs/iPSCs were efficiently induced to neurospheres by using
58 a newly established induction system in vitro. And It-NES cells derived from
59 hESCs/iPSCs-neurospheres using two induction system (coated vs AA)
60 both expressed neural pluripotency-associated genes PAX6, NESTIN,
61 SOX1, SOX2. After long-term cultivation, we found that they both can
62 maintain the long-term expansion for more than a dozen generations while
63 maintaining neuropluripotency. Moreover, the It-NES cells retain the ability
64 to differentiate into general functional neurons that highly express β -tubulin.
65 We also demonstrated that AA promotes the generation and long-term
66 expansion of It-NES cells by promoting collagen synthesis via the
67 MEK-ERK1/2 pathways.

68 **Conclusions:** Taken together, this new chemically defined culture system
69 is stable and effective to generate and culture the It-NES cells induced by
70 hESCs/iPSCs using serum-free medium combined with ascorbic acid (AA).
71 The It-NES cells under this culture system can maintain the long-term
72 expansion and neural pluripotency, with the potential to differentiate into
73 functional neurons.

74 **Keywords:** Spinal cord injury, Neurospheres, Ascorbic acid, It-NES cells,
75 Human pluripotent stem cells.

76

77 **Introduction**

78 Spinal cord injury (SCI) is a destructive neuro-degenerative disorder
79 with life-long consequences, and it often leads to irreversible changes due
80 to the loss of neurons and glial cells [1,2]. However, the quality of life for
81 SCI patients has been improved by the findings of past research, but there
82 is still no effective therapy available. In recent years, there are many
83 experimental animal models simulating SCI in humans and the
84 transplantation of neural stem/progenitor cells (NS/PCs) has been shown to
85 be an effective treatment to cure the neurological disorders and repair
86 damaged brain tissue due to their ability to differentiate into neurons and
87 glial cells and the competence to secrete neurotropic factors [3,4].

88 In the research field of SCI, NS/PCs derived from mouse embryonic
89 stem cells (mESCs) [5] and rat embryonic spinal cord[3] have been
90 transplanted into the injured spinal cord of mice and rats, promotes
91 functional recovery in animal SCI models. Also, the NS/PCs, derived from
92 the human fetal brain [6,7] have been transplanted into the injured spinal
93 cord model of mice or nonhuman primates [6,8], greatly promoted the

94 development of stem cell-based transplantation therapy for human patients.
95 However, the collection of NS/PCs from brains of aborted fetuses or surplus
96 embryos is not allowed on clinical applications in Japan. This fact has been
97 a major hinder to the clinical use of human NS/PCs[9].

98 Under this circumstance, human embryonic stem cells (hESCs) [10,11]
99 and induced pluripotent stem cells (iPSCs) established by Yamanaka et al.
100 [12,13] have the ability to differentiate into NS/PCs, providing a rich source
101 of neurons and glial cells [14], therefore, they provided a new approach for
102 studying human neurodevelopment, simulating neurological disease, and
103 regenerative medicine. Moreover, iPSCs reprogrammed from somatic cells
104 provide the new prospect of generating patient-specific cells for disease
105 modeling and regenerative medicine[12]. The ability of proliferation and
106 differentiation of iPSCs is almost equal to that of ES cells, and the use of
107 iPSCs could circumvent the ethical issues and rejection in NS/PCs
108 transplantation related to the use of embryos and aborted tissues.

109 Neurosphere is a well-known classic culture system for NS/PCs
110 extension [15] and the present neurospheres induction protocols of for
111 human pluripotent stem cells (hPSCs) is depended upon the formation of
112 embryoid body [16–18], but this method has many considerable
113 disadvantages, such as unclear culture conditions, prolonged differentiation
114 and low differentiation efficiency. In addition to these methods, human

115 long-term self-renewing neuroepithelial-like stem cells (It-NES cells) [18]
116 have reached the standard defined as NS/PCs, providing a more
117 homogeneous and robust cell generation in ways of monolayer adherent
118 cultures [19]. And all It-NES cells derived from different hESC/hiPSCs
119 exhibit characteristics like continuous expandability, stable neuronal and
120 glial differentiation competence and the capacity to generate functionally
121 mature human neurons [14,20,21]. However, these It-NES cells culture
122 systems are inseparable from the plates coated with adhesive materials, so
123 they are not chemically clear culture systems. Therefore, it is very crucial to
124 develop a culture system of It-NES cells under chemically defined
125 conditions.

126 In our research, we describe for the first time a serum-free culture
127 system to generate It-NES cells from different hPSCs under chemically
128 defined conditions without the need for pre-coated and avoiding the use of
129 basement membrane preparation such as Matrigel matrix,
130 poly-L-lysine/laminin-coated or inactivated MEF-feeder plates [22,23] with
131 unclear chemical composition, which limit the clinical application of NS/PCs.
132 In this system, we applied ascorbic acid (AA) which promoting collagen
133 synthesis through the MEK-ERK1/2 pathway [24] to induce the generation
134 and long-term expansion of It-NES cells. This method is expected to
135 facilitate the clinical application of NS/PCs derived from hPSCs for the

136 regenerative medicine of neurological disorders and injuries, like spinal
137 cord injury [25].

138 **Materials and Methods**

139 **1 Human PSCs culture**

140 Human PSCs including hESCs-H9 and urine-derived iPSCs (UiPSCs,
141 generated in our laboratory by standard Sendai Reprogramming
142 (CytoTune™-iPS 2.0 Sendai Reprogramming Kit, Invitrogen, USA)) were
143 cultured in PSCeasy medium (Cellapy, China) on six-well plates (Corning,
144 US) coated with a 1:500 dilution of hESC-matrigel (Corning). Medium was
145 changed every day. PSCs were passaged every 3–4 days at 70-80%
146 confluence with 0.5 mM EDTA (Cellapy). All cells were maintained at 37 °C,
147 5% CO₂ in incubator (Thermo Fisher Scientific, USA).

148 **2 Neurosphere differentiation of PSCs**

149 The protocol of neurosphere differentiation was shown in Figure 1A.
150 When hPSCs were grown for 4 days on human recombinant laminin
151 fragment iMatrix-511 (Nippi , Japan) - coated (0.5 µg/cm²) 6-well plate, at
152 which time they reached 100% confluence, namely at day 0 of induction of
153 differentiation, cells were washed with PBS (Hyclone, USA), and then
154 added the 2ml NeuroEasy Human Neural Stem Cell Induction Culture
155 medium (Cellapy, DMEM/F12 containing 500 µg/mL recombinant human

156 albumin (A0237, Sigma-Aldrich)+2 uM Chir99021(Selleck, China)) per well,
157 and change the medium every two days until the cells exhibited a rosette
158 morphology (5-6d). And then, cells were dissociated with Accutase
159 (Innovative Cell Technologies, USA) for 10min and resuspended by
160 NeuroEasy Human Neural Stem Cell Culture medium (Cellapy, DMEM/F12
161 containing 2% NeuroEasy Human Neural Stem Cell supplement (xeno-free
162 version of B27, Cellapy), supplemented with 20 ng/ml FGF2, 25ng/ml EGF
163 and 25ng/ml heprin) on low cell-adhesion plates. A ROCK inhibitor, Y27632
164 (Selleck, China) at a concentration of 10uM, was used only at the time of
165 plating. Next day, cell aggregated to form neurospheres, and the medium
166 was changed to Neural stem cell culture medium without Y27632.
167 Neurospheres were passaged every 3-4 days when they grow to 20um in
168 size with accutase. Medium was changed every 2-3days. All cells were
169 maintained at 37 °C, 5% CO₂ in incubator.

170 **3 It-NES cells differentiation of neurosphere**

171 When the neurospheres grow to 20um in size, we perform PBS
172 washing and dissociated neurospheres with Accutase for 10min and
173 resuspended by It-NES cells culture medium (DMEM/F12 containing 2% N2
174 (xeno-free version, Cellapy), supplemented with 20 ng/ml FGF2, 25ng/ml
175 EGF and 25ng/ml heprin) on gelatin-coated plate. When we identified the
176 effect of AA in the generation and expansion of It-NES, we use

177 uncoated-plate and we add the 50 µg/ml L-ascorbic acid-2-phosphate
178 (Selleck) in the It-NES cells culture medium during resuspension, long-term
179 culture and other experimental procedures.

180 **4 Cell viability and proliferation assays**

181 Cell viability and proliferation was measured by the Cell Counting Kit-8
182 (CCK-8) assay (Dojindo, Japan). The It-NES cells under different condition
183 were plated in 96-well plates (3000 cells/well) and cell viability was detected
184 after 12h according to the manufacturer's protocol. Cell proliferation was
185 detected every 24 h. At every point in time, 10 µl of CCK-8 reagent was
186 added to each well, and the plate was incubated for 2 h at 37 °C. And then,
187 each sample was measured absorbance at 450 nm using an automatic
188 microplate reader (BioTek Synergyn 4, USA). All experiments had three
189 technical replicates each.

190 **5 Differentiation of It-NES cells into generic neurons and glia**

191 It-NES cells were dissociated by for 10mins and plated on human
192 recombinant laminin fragment iMatrix-511-coated 12-well plate with glass
193 slides at a low density of 2×10^4 cells/cm² in Neuron maintenance medium
194 (DMEM/F12 containing 2% NeuroEasy Human Neural Stem Cell
195 supplement (xeno-free version of B27, Cellapy)), and half of the medium
196 was changed every 2 or 3 days.

197 **6 RNA extraction and quantitative real-time PCR**

198 Cells were lysed and harvested using TRIzol™ Reagent (Invitrogen,
199 USA) according to the manufacturer's instructions. RNA was quantified with
200 NANO drop 2000 (Thermo Fisher Scientific), and 1ug RNA was reverse
201 transcribed into cDNA using the PrimeScript™ RT Master Mix reverse
202 transcription System (TaKaRa, Japan). The levels of relative gene
203 expression were analyzed by quantitative reverse transcriptase PCR
204 (qRT-PCR) with TB Green™ Premix Ex Taq™ II (TaKaRa) using the
205 iCycler iQ5 (Bio-Rad, USA). The housekeeping gene GAPDH was used for
206 internal normalization and the relative quantification of gene expression
207 was calculated according to the Δ CT method. The qRT-PCR primers are
208 listed in Supplementary Table S1.

209 **7 RNASeq Processing and Biological Information Analysis**

210 Clustering and sequencing were executed by Novogene Corporation
211 (Tianjing, China). The clustering of the index-coded samples was
212 performed on a cBot Cluster Generation System using TruSeq PE Cluster
213 Kit v3-cBot-HS (Illumina) . After cluster generation, the library preparations
214 were sequenced on an Illumina Hiseq platform .Raw data (raw reads) of
215 fastq format were processed through in-house perl scripts. All the
216 downstream analyses were based on the clean data with high quality.

217 Differential expression analysis was performed using the DESeq2 R
218 package (1.16.1). The resulting P-values were adjusted using the
219 Benjamini and Hochberg's approach for controlling the false discovery
220 rate . Genes with an adjusted P-value <0.05 found by DESeq2 were
221 assigned as differentially expressed. Gene Ontology (GO) enrichment
222 analysis of differentially expressed genes was implemented by the
223 clusterProfiler R package, in which gene length bias was corrected. GO
224 terms with corrected Pvalue less than 0.05 were considered significantly
225 enriched by differential expressed genes. KEGG is a database resource for
226 understanding high-level functions and utilities of the biological system from
227 molecular-level information. We used clusterProfiler R package to test the
228 statistical enrichment of differential expression genes in KEGG pathways.

229 **8 Immunofluorescence staining**

230 Cells were fixed with 4% PFA (Solarbio, China) for 10-15min at room
231 temperature, permeabilized with 0.3% Triton X-100 (Sigma) for 10-15 min
232 at room temperature, blocked with 3% bovine serum albumin (Solarbio) for
233 45-60 min at room temperature, and then incubated with primary antibodies
234 against OCT4 (1:100; Santa cruz Biotechnology), SSEA4 (1:100; Santa
235 cruz Biotechnology), SOX2 (1:100; Santa cruz Biotechnology), NESTIN
236 (1:100; Sigma), β 3-tubulin (1:100; Abcam), and GFAP (1:100; Abcam)
237 overnight at 4 °C . And then cells were incubated with secondary antibodies:

238 Goat anti-Rabbit IgG Alexa Fluor 488 (1:200; Invitrogen) and Goat
239 anti-Mouse IgG Alexa Fluor 594 (1:200; Invitrogen) for 1 hour at 37 °C.
240 Wash with PBS three times before each step. Nuclei were stained with
241 DAPI (300 nM, Invitrogen) for 15 min at room temperature. Fluorescence
242 images were captured by Leica DMI 4000B fluorescence microscope and
243 Leica TCS SP5 MP confocal laser scanning microscope (Leica, Germany).

244 **9 Western blotting**

245 Cells were lysed using tissue protein extraction reagent (Thermo, USA)
246 containing phosphatase inhibitor cocktail (1:100, Thermo), protease inhibitor
247 cocktail (1:100, Thermo), and 5mM EDTA (Thermo), lysates were oscillated
248 and centrifuged (13 000 × g, 15 min, 4 °C). The supernatant was collected
249 and stored at -80 °C. The protein concentrations were determined using
250 the BCA protein assay kit (Thermo). The protein was mixed with 5X
251 SDS-PAGE protein loading buffer (Beyotime, China) and denatured by
252 100 °C water bath. Then, the samples denatured were electrophoresed in
253 10% SDS-PAGE and transferred to PVDF membranes using transfer
254 device (Bio-Rad). The membranes were blocked with 5% non-fat milk
255 prepared in TBST for 1 hour at 37 °C and then incubated at 4 °C overnight
256 with the primary antibodies: ERK1/2(1:1000, Cell Signaling Technology,
257 p44/42 MAPK Rabbit mAb, 4695), p-ERK1/2(1:1000, Cell Signaling
258 Technology, Phospho-p44/42 MAPK (Erk1/2) XP®, Rabbit mAb, 4370) and

259 the internal normalization mouse anti-GAPDH (1:1000, Santa cruz
260 Biotechnology). Next, the membranes were washed in TBST, incubated
261 with secondary antibody: Goat anti-Rabbit IgG (H + L) IRDye 800CW or
262 Goat anti-Rabbit IgG (H + L) IRDye 800CW (1:20 000; LI-COR) for 1 h at
263 37 °C. The images were observed with a UVA Bio Imaging System and
264 analyzed with ImageJ software.

265 **10 Data analysis and statistics**

266 All data are analyzed by means \pm standard errors of the means (S.E.M.).
267 Statistical significance was evaluated using two-sided t-test for two groups,
268 and using one-way ANOVA test for statistical differences of multiple groups.
269 Significant differences were considered when *P < 0.05, **P < 0.01.

270 **Results**

271 **Generation of neurosphere from hESCs/iPSCs.**

272 To evaluate the availability of our approach, studies were performed in
273 parallel in two cell lines, including hESC-H9 and hiPSC produced from urine
274 [26] cells by standard Sendai Reprogramming. Urine-iPSCs (UiPSs) are
275 morphologically identical from hESC-H9 and express representative
276 pluripotent markers, including SSEA4 and OCT4 (Figure.1B a,b,f,g).

277 Neurospheres for culturing NS/PCs in vitro have been reported and are
278 widely used [16], the protocol of neurospheres induction from

279 hESC/hiPSCs is shown in Figure 1A. hESC/hiPSCs can be efficiently
280 differentiated to neurospheres with the rosette formation using the neural
281 stem cell induction medium and rosette dissociation using the neural stem
282 cell medium. In the process of differentiation, the morphology of hPSCs
283 transition to neurospheres phenotypes is shown in Figure 1B. There is a
284 significant change in cell morphology along with the cell proliferation. At the
285 stages of hPSCs, cells expanded and displayed a round, colony
286 morphology (Figure.1B a,b,f,g). As the duration of neural induction to day 5,
287 cells exhibited a rosette morphology (Figure.1B c,h). At day 6, cells were
288 digested and resuspended to form neurospheres (Figure.1B d,i). By starting
289 with 100% confluence pluripotent stem cells, it is possible to get 20–40-fold
290 increases in cell number of NS/PCs in 6 days, so we can get enough NSCs
291 to meet the needs of clinical applications.

292 Gene relative expression analysis shown in Figure 1C confirmed the
293 identity of neurospheres at day7 through the progressive loss of hPSCs
294 pluripotency markers such as *NANOG* and *OCT4*, and a significantly
295 increase in the expression of neural progenitor markers *SOX1*, *PAX6*,
296 *NESTIN* and *SOX2* which defining early neural fate. Neurospheres at day7
297 of neural induction were plated for immunostaining with pluripotent NS/PCs
298 markers and most of the cells were positive for NS/PCs markers *SOX2* and
299 *NESTIN* (Figure.1B e,j). Neurospheres generation and gene expression

300 was evaluated using two cell lines H9 and UiPS, and no significant
301 difference was observed (Figure.1C&Figure.1D).

302 These results suggest that neurospheres derived from hESC/hiPSCs
303 by neural induction medium possess the NS/PCs phenotype.

304 **AA promoted the generation of It-NES cells from neurospheres under**
305 **chemically defined conditions.**

306 Whereas neurospheres exhibit heterogeneous character and tend to
307 reproduce the neural development process, monolayer adherent cultures
308 provide a more robust and homogeneous cell generation [19].

309 We have tried many ways to get It-NES cells (data not shown), and we
310 found that ascorbic acid (AA) can promote the attachment of It-NES cells.
311 To evaluate the efficiency of using AA to induce It-NES cells, we
312 subsequently performed paralleled derivation of It-NES cells from
313 H9/UiPS-neurospheres using conventional N2 medium with gelatin-coated
314 and N2+AA medium without gelatin-coated. The It-NES cell derivation
315 protocol from neurospheres is shown in Figure 2A. Neurospheres were
316 dissociated into single cells and plated on gelatin-coated/uncoated plates in
317 the N2 medium with FGF2 and EGF (Coated/Uncoated) or plated on
318 uncoated plates in the N2+AA medium with FGF2 and EGF (AA). The
319 morphologies of the It-NES cells derived from H9/UiPS-neurospheres in N2
320 medium (coated) and N2+AA medium (AA) were comparable with each

321 other (Figure.2B a,c vs. d,f), and there were few It-NES cells in uncoated
322 plate (Figure.2B b,e). The viability of It-NES cells in different culture
323 condition was evaluated by a CCK-8 assay (Figure.2C), and there were no
324 differences in the cell viability of the It-NES cells in coated and AA
325 conditions after plated for 12h. Growth kinetics showed no significant
326 difference in proliferation of the It-NES cells in coated and AA conditions
327 (Figure.2D), and we can see from this result that It-NES cells expanded in
328 N2+AA medium (AA) generate large numbers of cells, almost two-fold
329 expansion can be achieved upon each passage, and It-NES cells were
330 passaged once a week on average.

331 All cells cultured in two methods in passage one expressed NS/PCs
332 markers *SOX1*, *PAX6*, *NESTIN* and *SOX2* (Figure.3A&3B), and had low
333 expression of hPSCs pluripotency markers *NANOG* and *OCT4*. What
334 deserves our attention is that N2+AA (AA) cultured cells have higher
335 expression levels of NS/PCs markers than N2 cultured (coated) cells, which
336 indicates that the AA maybe can promote the neural pluripotency of It-NES
337 cells. The immunocytochemical analysis indicated that there were no
338 significant differences in the percentage of NS/PCs marker-positive cells
339 (Figure.3C, 3D&3E), however, there are slightly more SOX2-positive cells
340 in N2+AA culture, demonstrating the optimal culture using N2+AA.

341 These data suggest that N2+AA without gelatin-coated is suitable for
342 the induction and growth of It-NES cells.

343

344 **AA can maintain the long-term expansion and neural pluripotency of**
345 **It-NES cells**

346 To verify whether the N2+AA culture system supports excellent
347 proliferation and preserves the neural pluripotency of It-NES cells, we
348 cultured the cells of N2 cultured (coated) and N2+AA cultured (AA) to 15
349 generations and identified the capacity of proliferative and the potential of
350 neural differentiation by qPCR and immunochemistry. Gene relative
351 expression analysis by qPCR shown in Figure 4A&4B displayed that cells
352 cultured in N2 (coated) or N2+AA (AA) in passage 5 and passage 10
353 expressed NSC markers *SOX1*, *PAX6*, *NESTIN* and *SOX2*, and had almost
354 no expression of pluripotency markers *NANOG* and *OCT4*. And analyses of
355 expanded It-NES cells by immunofluorescence staining showed consistent
356 expression of NSC markers, including *SOX2* (> 75%) and *NESTIN*
357 (>95%) at passage 5 and passage 10 (Figure.4C&D). Furthermore, N2+AA
358 can maintain normal karyotype over 15 passages (Figure.4E).

359 The above results reveal that N2+AA without gelatin-coated can
360 maintain the long-term expansion and excellent proliferation, and preserve
361 the neural pluripotency of It-NES cells.

362 **It-NES cells cultured in N2+AA can differentiate into functional**
363 **neurons**

364 The neurons generated from NSCs provide a useful model for studying
365 human disease, drug screening, toxicity testing, and cell therapy. To test
366 the neuronal differentiation potential of It-NES cells cultured by N2+AA
367 medium, we assessed whether the It-NES cells could differentiate into
368 generic neurons. As shown in Figure 5A a,c, after two weeks of neuronal
369 differentiation, H9-It-NES cells in N2 cultured (coated) and N2 AA (AA)
370 culture systems can differentiate into a large number of neurons. And this
371 phenomenon can be repeated by UiPS-It-NES cells, they could also
372 differentiated into substantial numbers of neurons (Figure.5B).As
373 demonstrated by the immunocytochemical of the neuronal marker class III
374 β -tubulin and the glia marker glial fibrillary acidic protein (GFAP) (Figure.5A
375 b,d) respectively, the neurons produced by our system are highly pure (\geq
376 95% class III β -tubulin-positive neurons; < 5% GFAP-positive glia) and can
377 be maintained in long-term culture. Quantitative RT-PCR analysis shows
378 that neurons generated by neuronal differentiation of N2 cultured (coated)
379 and N2+AA (AA) cultured It-NES cells exhibit similar gene expression
380 patterns (Figure.5C): after 14 days in neuron culture medium, the
381 expression of neuronal type markers *NeuN*, *Neurog2* and *β -tubulin*, and
382 glutamatergic neuron markers *vGlut1* was up-regulated, and the expression

383 of the astrocyte markers *S100B-β* and *GFAP* was low. Subsequently, we
384 investigated the electrophysiological activity of neurons differentiated from
385 H9-It-NES cells using a microelectrode array (MEA) recording system. We
386 plated It-NES cell-derived neurons on the MEA plate, then we cultured them
387 in the Neuron maintenance medium and recorded the spontaneous activity
388 (Figure 5D). The spontaneous activity of the It-NES cell-derived neurons
389 gradually increased and maintained a high level for 50 days after
390 differentiation (Figure 5E). This indicate that N2+AA cultured It-NES
391 cell-derived neurons are functional and able to maintain long-term
392 spontaneous neuronal activity.

393 These data indicate that N2+AA cultured It-NES cells have the capacity
394 to differentiate into generic functional neurons.

395 **Transcriptome analysis of It-NES cells**

396 In order to explore the changes in gene expression, functional
397 consequences, and the potential molecular mechanism caused by ascorbic
398 acid, we performed RNA-seq analysis on N2 cultured (coated) and N2+AA
399 (AA) cultured It-NES cells. We found that 3050 genes were significantly
400 upregulated and 1967 downregulated after AA cultured (Figure 6A).

401 In order to understand the biological mechanism of AA effects, we
402 performed GO (Gene ontology) enrichment analysis on differentially

403 expressed genes (DEG). The 10 most significant categories were identified
404 in the cell composition, molecular function and biological process groups
405 respectively (Figure. 6B). Most remarkable categories are concentrated in
406 regulation of extracellular matrix (ECM) assembly, collagen metabolic
407 process within the biological process, and extracellular matrix component
408 within the cell composition (Figure. 6B). Among them, cell adhesion
409 molecule binding was the most different categories (Figure. 6C). The above
410 results indicated that AA plays a critical role in cell adhesion, ECM
411 deposition/remodeling and collagen synthesis, which is consistent with
412 previous researches [27,28]. To understand the biological impact of these
413 differentially expressed transcripts, we next performed KEGG (Kyoto
414 Encyclopedia of Genes and Genomes) functional analysis (Figure. 6D).
415 Among 20 significantly different pathways, these pathways were mainly
416 related to cell adhesion and ECM-receptor interaction.

417 Overall, these transcriptional profiling results indicate that AA is
418 involved in the regulation of cell adhesion and is closely related to the
419 regulation of ECM deposition/remodeling, especially in collagen synthesis.

420 **AA promotes the generation of It-NES cells through the MEK-ERK1/2**
421 **pathway by promoting collagen synthesis.**

422 Transcriptome analysis indicates that ascorbic acid plays a role by
423 affecting cell adhesion, ECM deposition/remodeling and collagen synthesis.
424 However, it is known that ascorbic acid is also an antioxidant, so we need
425 to determine whether the generation of It-NES cells by AA is due to its
426 antioxidant properties. We found that treatment with other antioxidants such
427 as vitamin B1 (Vb1) and reduced glutathione (GMEE) could not mimic the
428 effect of AA on It-NES cell adhesion (Figure 7A), indicating that the effect of
429 AA on promoting adherence of It-NES cells is not related to its antioxidant
430 properties. Collagens are indispensable components of ECM, which play a
431 vital role in cell development and function, and have been shown to affect
432 cell proliferation and differentiation [29]. In addition, It has been pointed
433 out that AA enhanced the proliferation of CPCs via the MEK-ERK1/2
434 pathway through manipulating collagen synthesis [24]. In order to verify
435 whether the mechanism of AA promoting the generation and expansion of
436 It-NES cells is in the same way as the above article, we analyzed the effect
437 of AA on collagen synthesis and found that the expression of collagen
438 genes *Col1a1* and *Col4a1* of It-NES cells cultured by N2+AA (AA)
439 increased significantly (Figure.7B), which is consistent with the
440 transcriptome results. In order to further explore the role of AA in the
441 synthesis of collagen, we used a generic collagen synthesis inhibitor
442 L-2-Azetidine carboxylic acid (AzC), and we found that the adhesion of
443 It-NES cells by AA were completely eliminated by AzC (50 $\mu\text{mol/l}$)

444 (Figure.7C). And then, to confirm whether AA promotes collagen synthesis
445 via the MEK-ERK1/2 pathway, we performed the validation of AA-induced
446 ERK1/2 activation and the effect of ERK1/2 inhibitors. Western blot analysis
447 shown in Figure 7D&7E indicated that the level of p-ERK1/2 was higher in
448 N2+AA (AA) cultured It-NES cells than N2 cultured (coated). Moreover, the
449 ERK1/2 inhibitor (LY3214996) completely abolished AA-induced generation
450 and expansion of N2+AA (AA) cultured It-NES cells (Figure.7F), whereas
451 the N2 cultured remained unchanged when addition of LY3214996,
452 suggesting that the MEK-ERK1/2 pathway is involved in the AA-dependent
453 generation and expansion of It-NES cells.

454 The above data demonstrate that the AA-induced collagens synthesis
455 promotes the generation and expansion of It-NES cells through the
456 MEK-ERK1/2 pathway.

457 **Discussion**

458 In this study, a new system was established to induce It-NES cells from
459 hESC/hiPSCs-neurospheres under chemically defined conditions and this
460 system designed to meet the requirements of quality in clinical settings.
461 This system also demonstrated that N2+AA culture system is suitable for
462 the induction and long-term expansion of It-NES cell (Figures 2,3,4), and
463 the differentiation of generic neurons (Figure 5). AA added in N2 medium

464 for generation and expansion of It-NES cells avoids pre-coating and avoids
465 the use of basement membrane preparations, therefore, the chemical
466 composition of this culture system is more specific and this culture system
467 is subject to the standards of serum-free system for clinical application in
468 the future. Moreover, this system is suitable for various pluripotent stem
469 cells, including ESCs and iPSCs, therefore, we can use the patient-specific
470 iPSCs to avoid ethical issues and immune rejection.

471 Neural stem/progenitor cells (NS/PCs) can maintain long-term
472 expansion, constantly self-renew, and have the ability to differentiate into a
473 variety of neural cell lineages, including neurons and astrocytes, which
474 illustrating the potential of neural stem cells in cell replacement therapies of
475 neurological disorders and injuries caused by the loss of neurons and glial
476 cells, such as spinal cord injury [30,31]. Neurospheres, a classic culture
477 system of expanding NS/PCs, it's cultivation as a free-floating aggregate
478 and it is considered as a more natural environment for cells due to the 3D
479 niche-like structure [15]. Methods of neurospheres induction and expansion
480 are slightly changed from the previous protocols. Neurospheres are usually
481 derived from single-cell suspensions of neural stem and progenitor cells
482 isolated from adult or fetal central nervous system (CNS), but neurospheres
483 can also be established from ES cells [32]. In our research, we used neural
484 stem cell induction medium to induce hESC/hiPSCs into rosettes, and after

485 the rosette was digested, they were spread on uncoated plate in the form of
486 a single cell suspension, then the single cells aggregate to form a spherical
487 shape and expand continuously. (Figure 1). The medium was
488 supplemented with B27 containing fibroblast growth factor (bFGF) and
489 epidermal growth factor (EGF) [33]. This neurosphere induction method is
490 proposed for the first time in this study, and it has a higher yield and a
491 higher efficiency than previous methods.

492 However, heterogeneity is one of the significant limitations of
493 neurospheres culture system [34]. This heterogeneity is because the size of
494 the neurospheres is difficult to control and usually not uniform, and cells
495 inside and outside the neurospheres are exposed to different environments.
496 The heterogeneity makes it difficult to maintain long-term expansion and
497 excellent differentiation potential. Moreover, after transplantation of
498 neurospheres into the nervous system, the neurogenic potential is gradual
499 loss and the yield of neurons decreases [34]. Therefore, there is a demand
500 to develop a new NS/PCs culture system that should be homogeneous,
501 robust, stable over time, and can constantly produce a large number of
502 neurons. Culture of NSCs in monolayer has been studied previously to
503 solve above mentioned needs, and human long-term self-renewing
504 neuroepithelial-like stem cells (lt-NES cells) are NS/PCs in a monolayer
505 culture system. They are a fairly homogeneous population of

506 undifferentiated cells with the ability to continuously expand and stably
507 produce functional neurons and glial [21]. The most common method to
508 generate It-NES cells is to pre-coat the culture plates with basement
509 membrane preparations rich in extracellular matrix (ECM), like laminin [35],
510 fibronectin, gelatin [36], and Matrigel [37]. However, basement membrane
511 preparations listed above all contain exogenous components, and these
512 exogenous components hinder the progress of cell transplantation due to
513 uncertain chemical composition. In our research, we have established a
514 new system for inducing and maintaining It-NES cells under chemically
515 defined conditions by using ascorbic acid (AA) (Figure.2A). The It-NES cells
516 cultured under this new system can stably expand in large numbers and
517 preserve the neural pluripotency (Figure.2-3), and can maintain long-term
518 culture (Figure.4). Moreover, It-NES cells cultured under our system can
519 maintain the differentiation potential of functional neurons (Figure.5).
520 Therefore, it is expected that this system will contribute to regenerative
521 medicine for spinal cord injury and other central nervous system (CNS)
522 diseases in the future, because this method can reliably provide a large
523 number of functional cells for regenerative medicine.

524 Ascorbic acid (AA) is widely known as an essential nutrient for guinea
525 pigs and primates [38,39]. Ascorbic acid has two major biological activities,
526 it serves as an antioxidant, and a cofactor of collagen synthesis [40]. In our

527 study, other antioxidants were tested, such as vitamin B1 (Vb1) and
528 reduced glutathione (GMEE), but none of them mimicked the effect of AA
529 on It-NES cells (Figure.7A), suggesting that antioxidant activity is not a
530 critical factor in the generation and expansion of It-NES cells. Therefore, we
531 focus on the role of ascorbic acid in the synthesis of collagen, as the
532 extracellular matrix (ECM) may affect cell adhesion. Ascorbic acid (AA) is
533 essential for the biosynthesis of collagen. It is a cofactor of prolyl and lysyl
534 hydroxylase and a stimulator of collagen gene expression. After addition of
535 AA to the culture medium, many studies have evaluated the relationship
536 between AA and collagen expression on the short- and long-term effects of
537 cells [27]. This study evaluated the use of AA in the long-term culture of
538 It-NES cells, and from the RNAseq results (Figure.6), it can be seen that AA
539 affects cell acquisition and long-term culture by affecting cell adhesion,
540 ECM remodeling and collagen synthesis. Moreover, during the long-term
541 culture, the expression of collagen genes *Col1a1* and *Col4a1* was
542 enhanced (Figure.7B), and we used one general collagen synthesis
543 inhibitor, L-2-Azetidine carboxylic acid (AzC) and found that the adhesion of
544 It-NES cells by AA were completely eliminated by AzC (50 $\mu\text{mol/l}$)
545 (Figure.8C). These data indicate that the AA plays a promotive role in the
546 generation and expansion of It-NES cells is due to collagen synthesis rather
547 than antioxidant properties. It has been reported that AA specifically
548 enhances CPC proliferation by manipulating collagen synthesis through the

549 MEK-ERK1 / 2 pathway [24], consistent with our results, the MEK-ERK1/2
550 pathway is activated by AA and the adhesion of It-NES cells by AA were
551 completely abolished by ERK inhibitor (1 μ M) (Figure 7D-F). Moreover, we
552 found that It-NES cells cultured in N2+AA (AA) have higher expression
553 levels of NS/PCs markers than cells cultured in N2 (coated) (Figure 2-3).
554 The above studies indicate that AA may play a critical role in development
555 of NS/PCs, and may provide a more suitable culture environment for It-NES
556 cells and promote the neural pluripotency of It-NES cells.

557 **Conclusion**

558 We have successfully established a new culture system for the
559 long-term large-scale monolayer culture of NS/PCs from hESC/hiPSCs. AA
560 promotes a homogenous population of It-NES cells grown in an
561 environment without foreign components. After 10-15 passages, the cells
562 on the N2+AA can maintain good self-renewal ability, and the cells can
563 maintain neural pluripotency and the ability to differentiate to generic
564 functional neurons. In addition, we have demonstrated that AA specifically
565 promotes the generation and expansion of neuroepithelial-like stem cells
566 through the MEK-ERK1 / 2 pathway by increasing collagen synthesis.
567 Findings from this study help promote the clinical application of
568 hPSC-derived NS/PCs in regenerative medicine for spinal cord injury and
569 neurological diseases.

570 **Abbreviations**

571 SCI: Spinal cord injury; hPSCs: Human pluripotent stem cells; NS/PCs:
572 neural stem/progenitor cells; AA: ascorbic acid; It-NES cells: long-term
573 self-renewing neuroepithelial-like stem cells; RT-PCR: reverse
574 transcription-polymerase chain reaction; MEA: microelectrode array;
575 mESCs: mouse embryonic stem cells; hESCs: human embryonic stem cells;
576 iPSCs: induced pluripotent stem cells; UiPSs: Urine-iPSCs; GFAP: glial
577 fibrillary acidic protein; ECM: extracellular matrix; GO: Gene ontology;
578 KEGG: Kyoto Encyclopedia of Genes and Genomes; Vb1: vitamin B1;
579 GMEE: reduced glutathione; AzC: L-2-Azetidine carboxylic acid; CNS:
580 central nervous system; bFGF: fibroblast growth factor; EGF: epidermal
581 growth factor.

582 **Declarations**

583 **Ethics approval and consent to participate**

584 Not applicable

585 **Consent for publication**

586 Not applicable

587 **Availability of data and materials**

588 All data generated or analysed during this study are included in this
589 published article.

590 **Competing interests**

591 The authors declare that they have no competing interests.

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596 **Authors' contributions**

597 LF and LWJ conceived the idea and designed the experiments, BR and CY
598 performed cell experiments, data analysis. BR and Amina performed
599 manuscript preparation. WFJ, TL, and ZSY are responsible for cell-culture
600 experiments and the collection and assembly of data. LYN, MSH,DT
601 contributed to molecular experiments. GTW and YY contributed to function
602 analysis. All authors read and approved the final manuscript.

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757

758

759 **FIGURE LEGENDS**

760 **Figure 1. Generation of neurosphere from hiPS/ESCs**

761 (A)The protocol for deriving neurospheres from H9 and UiPS.(B) The
762 morphology and Immunostaining of H9 and UiPS at days -1 (a,b,f,g), day 5

763 (c,h), and day 7 (d,e,i,j) during neural induction. Immunostaining showing
764 emergence of SSEA4+ (green), OCT4+(red) H9 and UiPS(b,g), and
765 SOX2+ (red),NESTIN+ (green) neurospheres induced from H9 and UiPS at
766 day 7(e,j). Scale bars = 100 μ m. Nuclei were counterstained with DAPI
767 (blue). (C) Quantitative RT-PCR analysis of the expression level of neural
768 progenitor markers *SOX1*, *PAX6*, *NESTIN* and *SOX2*, and pluripotency
769 markers *NANOG* and *OCT4* of H9 and H9-neurospheres. (D) Quantitative
770 RT-PCR analysis of the expression level of neural progenitor markers
771 *SOX1*, *PAX6*, *NESTIN* and *SOX2*, and pluripotency markers *NANOG* and
772 *OCT4* of UiPS and UiPS-neurospheres. n = 3 each. Data are expressed as
773 means \pm SD.*P < 0.05, **P < 0.01 vs. control.

774

775 **Figure 2. It-NES cells can be generated from H9/UiPS-neurospheres**
776 **under a chemically defined culture system using N2+AA medium.**

777 (A)The It-NES cells derivation protocol from neurospheres. (B) The
778 morphologies of the It-NES cells in N2 medium (coated/uncoated) and
779 N2+AA (AA). Scale bars = 100 μ m. (C) The viability of P0 It-NES cells was
780 assessed by a CCK-8 assay after plated for 12 h. (D) Expansion capacity of
781 It-NES cells investigated by continuous culture in N2 medium
782 (coated/uncoated) and N2+AA (AA) in 8 days. n = 3 each.

783

784 **Figure 3. N2+AA medium can maintain the neural pluripotency of**
785 **It-NES cells**

786 (A) Quantitative RT-PCR analysis of the expression level of NSC markers
787 *SOX1*, *PAX6*, *NESTIN* and *SOX2*, and pluripotency markers *NANOG* and
788 *OCT4* of P0 H9 neurospheres-derived It-NES cells. (B) Quantitative
789 RT-PCR analysis of the expression level of NSC markers *SOX1*, *PAX6*,
790 *NESTIN* and *SOX2*, and pluripotency markers *NANOG* and *OCT4* of P0
791 UiPS neurospheres-derived It-NES cells. n = 3 each. Data are expressed
792 as means \pm SD.*P < 0.05, **P < 0.01 vs. control. (C) Immunofluorescence
793 for NSC markers SOX2 (red) and NESTIN (green) of P0 H9 & UiPS
794 neurospheres-derived It-NES cells. Scale bars = 100 μ m. Nuclei were
795 counterstained with DAPI (blue). (D) Quantification of SOX2, NESTIN
796 positive cells of Immunofluorescence of P0 H9 neurospheres-derived
797 It-NES cells. P, passage.(E) Quantification of SOX2, NESTIN positive cells
798 of Immunofluorescence of P0 UiPS neurospheres-derived It-NES cells.

799

800 **Figure 4. It-NES cells can maintain long-term expansion and neural**
801 **pluripotency using N2+AA medium.**

802 (A) Quantitative RT-PCR analysis of the expression level of NSC markers
803 *SOX1*, *PAX6*, *NESTIN* and *SOX2*, and pluripotency markers *NANOG* and
804 *OCT4* of P5 H9 neurospheres-derived It-NES cells. (B) Quantitative
805 RT-PCR analysis of the expression level of NSC markers *SOX1*, *PAX6*,

806 *NESTIN* and *SOX2*, and pluripotency markers *NANOG* and *OCT4* of P10
807 H9 neurospheres-derived It-NES cells. n = 3 each. Data are expressed as
808 means \pm SD.*P < 0.05, **P < 0.01 vs. control. (C) Immunofluorescence for
809 NSC markers *SOX2* (red) and *NESTIN* (green) of P5&P10 H9
810 neurospheres-derived It-NES cells. Scale bars = 100 μ m. Nuclei were
811 counterstained with DAPI (blue). (D) Quantification of *SOX2*⁺, *NESTIN*⁺
812 cells of Immunofluorescence. P, passage. (E) Representative karyotypes of
813 H9 neurospheres-derived It-NES cells at passage 15.

814

815 **Figure 5. It-NES cells differentiated into functional neurons.**

816 (A) Neurons and glia differentiated from P10 H9 neurospheres-derived
817 It-NES cells (a, c). Immunofluorescence for neuronal marker class III
818 β -tubulin (green) and glia marker GFAP (red) of Neurons and glia (b, d). (B)
819 Neurons and glia differentiated of P10 UiPS neurospheres-derived It-NES
820 cells (a, c). Immunofluorescence for neurons marker class III β -tubulin
821 (green) and glia marker GFAP (red) of Neurons and glia (b, d). Scale bars =
822 100 μ m. Nuclei were counterstained with DAPI (blue).

823 (C) Quantitative RT-PCR analysis of the expression level of neurons
824 markers NeuN, β -tubulin, vGlut1, and Neurog, and glia markers S100B- β
825 and GFAP . Transcript levels were normalized to undifferentiated It-NES
826 cells ,n = 3 each. (D) Representative images of activity map of neurons
827 differentiated from H9 neurospheres-derived It-NES cells on the

828 24-electrode array. (E) Spikes of neurons differentiated from H9
829 neurospheres-derived It-NES cells in 30s.

830

831 **Figure 6. Transcriptome analysis of It-NES cells cultured by coated or**
832 **AA .**

833 (A) Volcano plot of upregulated (red) or downregulated (green) transcripts
834 in AA cultured It-NES cells vs coated cultured. (B) Significantly enriched
835 Gene Ontology (GO) terms in AA cultured It-NES cells.

836 (C) GO terms categorization and distribution of differentially expressed
837 genes. GO terms were processed and categorized under three main
838 categories (biological process, cellular component, and molecular function).

839 (D) KEGG (Kyoto Encyclopedia Genes and Genomes) pathway rich detail.

840

841 **Figure 7. AA activates ERK signaling in a collagen**
842 **synthesis-dependent manner.**

843 (A) The morphology of It-NES cells treated with AA or alternative
844 antioxidants vitamin B1(Vb1) and reduced glutathione(GMEE). (B)

845 Quantitative RT-PCR analysis of the relative expression of collagen genes
846 *Col1a1* and *Col4a1*. n = 3 each. Data are expressed as means \pm SD.*P <

847 0.05, **P < 0.01 vs. control. (C) The morphology of It-NES cells treated with

848 AA and collagen synthesis inhibitor AzC. (D) The whole-cell extracts of

849 N2 cultured (coated) and N2+AA cultured (AA) It-NES cells were analyzed

850 by Western blot with phospho-specific antibodies against ERK1/2
 851 (pERK1/2), and total ERK1/2. GAPDH was used as a control. (E) ERK
 852 activation is defined as the ratio of p-ERK/total ERK. n = 3 each. Data are
 853 expressed as means \pm SD.*P < 0.05, **P < 0.01 vs. control. (F) The
 854 morphology of It-NES cells when treated with or without ERK inhibitor.
 855 Scale bars = 100 μ m.

856

857 **Supplementary**

858 Table S1. Primer sequences used for q-PCR

Gene	Forward 5'-3'	Reverse 3'-5'
PAX6	CTGAGGAATCAGAGAAGACAGGC	ATGGAGCCAGATGTGAAGGAGG
NESTIN	TCAAGATGTCCCTCAGCCTGGA	AAGCTGAGGGAAGTCTTGGAGC
SOX1	GAGTGGAAGGTCATGTCCGAGG	CCTTCTTGAGCAGCGTCTTGGT
SOX2	GCTACAGCATGATGCAGGACCA	TCTGCGAGCTGGTCATGGAGTT
OCT4	CCTGAAGCAGAAGAGGATCACC	AAAGCGGCAGATGGTCGTTTGG
NANOG	CTCCAACATCCTGAACCTCAGC	CGTCACACCATTGCTATTCTTCG
NeuN	TACGCAGCCTACAGATACGCTC	TGGTTCCAATGCTGTAGGTCGC

TUBB3	TCAGCGTCTACTACAACGAGGC	GCCTGAAGAGATGTCCAAAGGC
NEURO G	CAAGCTCACCAAGATCGAGACC	AGCAACACTGCCTCGGAGAAGA
vGlu1	GCAAGTACATCGAGGACGCCAT	GCCACGATGATGGCATAGACTG
GFAP	CTGGAGAGGAAGATTGAGTCGC	ACGTCAAGCTCCACATGGACCT
S100B	ACGTCAAGCTCCACATGGACCT	TCCTGGAAGTCACATTCGCCGT
Col1a1	GATTCCCTGGACCTAAAGGTGC	AGCCTCTCCATCTTTGCCAGCA
Col4a1	TGTTGACGGCTTACCTGGAGAC	GGTAGACCAACTCCAGGCTCTC
GAPDH	GGAGCGAGATCCCTCCAAAAT	GGCTGTTGTCATACTTCTCATGG

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