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| **NL-VS-L0 GO Enrichment (Cellular Component)** |
| **#** | **GO ID** | **Description** | **GeneRatio (8)** | **BgRatio (1849)** | **pvalue** | **p.adjust** |

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| 1 | [GO:0005875](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene1) | microtubule associated complex | 1 (12.5%) | 23 (1.24%) | 0.095460 | 0.731758 |
| 2 | [GO:0044430](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene2) | cytoskeletal part | 1 (12.5%) | 31 (1.68%) | 0.126741 | 0.731758 |
| 3 | [GO:0009536](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene3) | plastid | 3 (37.5%) | 330 (17.85%) | 0.157346 | 0.731758 |
| 4 | [GO:0015630](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene4) | microtubule cytoskeleton | 1 (12.5%) | 53 (2.87%) | 0.207935 | 0.731758 |
| 5 | [GO:0005856](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene5) | cytoskeleton | 1 (12.5%) | 63 (3.41%) | 0.242603 | 0.731758 |
| 6 | [GO:0031976](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene6) | plastid thylakoid | 1 (12.5%) | 65 (3.52%) | 0.249375 | 0.731758 |
| 7 | [GO:0031984](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene7) | organelle subcompartment | 1 (12.5%) | 66 (3.57%) | 0.252741 | 0.731758 |
| 8 | [GO:0030312](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene8) | external encapsulating structure | 1 (12.5%) | 70 (3.79%) | 0.266073 | 0.731758 |
| 9 | [GO:0071944](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene9) | cell periphery | 1 (12.5%) | 78 (4.22%) | 0.292115 | 0.731758 |
| 10 | [GO:0005623](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene10) | cell | 7 (87.5%) | 1353 (73.17%) | 0.322734 | 0.731758 |
| 11 | [GO:0044464](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene11) | cell part | 7 (87.5%) | 1353 (73.17%) | 0.322734 | 0.731758 |
| 12 | [GO:0009579](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene12) | thylakoid | 1 (12.5%) | 96 (5.19%) | 0.347773 | 0.731758 |
| 13 | [GO:0043229](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene13) | intracellular organelle | 5 (62.5%) | 971 (52.51%) | 0.419333 | 0.731758 |
| 14 | [GO:0043226](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene14) | organelle | 5 (62.5%) | 973 (52.62%) | 0.421806 | 0.731758 |
| 15 | [GO:0044424](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene15) | intracellular part | 6 (75%) | 1266 (68.47%) | 0.512928 | 0.731758 |
| 16 | [GO:0043234](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene16) | protein complex | 1 (12.5%) | 162 (8.76%) | 0.520496 | 0.731758 |
| 17 | [GO:0044435](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene17) | plastid part | 1 (12.5%) | 162 (8.76%) | 0.520496 | 0.731758 |
| 18 | [GO:0043231](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene18) | intracellular membrane-bounded organelle | 4 (50%) | 838 (45.32%) | 0.530658 | 0.731758 |
| 19 | [GO:0005622](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene19) | intracellular | 6 (75%) | 1282 (69.33%) | 0.534799 | 0.731758 |
| 20 | [GO:0043227](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene20) | membrane-bounded organelle | 4 (50%) | 854 (46.19%) | 0.550789 | 0.731758 |
| 21 | [GO:0044444](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene21) | cytoplasmic part | 3 (37.5%) | 658 (35.59%) | 0.586429 | 0.731758 |
| 22 | [GO:0005737](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene22) | cytoplasm | 3 (37.5%) | 661 (35.75%) | 0.590257 | 0.731758 |
| 23 | [GO:0044422](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene23) | organelle part | 2 (25%) | 435 (23.53%) | 0.595593 | 0.731758 |
| 24 | [GO:0044446](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene24) | intracellular organelle part | 2 (25%) | 435 (23.53%) | 0.595593 | 0.731758 |
| 25 | [GO:0043228](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene25) | non-membrane-bounded organelle | 1 (12.5%) | 218 (11.79%) | 0.634191 | 0.731758 |
| 26 | [GO:0043232](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene26) | intracellular non-membrane-bounded organelle | 1 (12.5%) | 218 (11.79%) | 0.634191 | 0.731758 |
| 27 | [GO:0032991](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene27) | macromolecular complex | 1 (12.5%) | 417 (22.55%) | 0.871137 | 0.934133 |
| 28 | [GO:0031224](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene28) | intrinsic component of membrane | 1 (12.5%) | 418 (22.61%) | 0.871857 | 0.934133 |
| 29 | [GO:0044425](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene29) | membrane part | 1 (12.5%) | 503 (27.2%) | 0.921585 | 0.953364 |
| 30 | [GO:0016020](file:///E%3A%5C2018-7-3%E7%81%AB%E9%BE%99%E6%9E%9C%E8%BD%AC%E5%BD%95%E7%BB%84%E6%B5%8B%E5%BA%8F%5C%E5%AE%8C%E6%95%B4%E7%89%88%E6%95%B0%E6%8D%AE%5CGDR3855-Hylocereus_undulatus_Britt-12-RNAseq_result%5C4_Function%5C2_Group_Diff_Function%5CUP_DOWN%5CGO%5CNL-VS-L0.C.html#gene30) | membrane | 1 (12.5%) | 770 (41.64%) | 0.986697 | 0.986697 |