P01 – BLAST - Hypericum androsaemum (texto texto)

general

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| Description | Scientific Name | Max Score | Total Score | Query Cover | E value | Per. Ident | Acc. Len | Accession |
| Prunus dulcis chloroplast, complete genome | Prunus dulcis | 1592.93 | 1592.93 | 99.77 | 0.0 | 100.0 | 157870 | KY101152.1 |
| Prunus dulcis DNA, unplaced-scaffold\_131 | Prunus dulcis | 1592.93 | 1592.93 | 99.77 | 0.0 | 100.0 | 40658 | AP020468.1 |
| Prunus dulcis DNA, pseudomolecule Pd02 | Prunus dulcis | 1592.93 | 3185.86 | 99.77 | 0.0 | 100.0 | 24875702 | AP019298.1 |
| Prunus dulcis plastid, complete genome >gi|1184802381|gb|KY085904.1| Prunus dulcis plastid, complete genome | Prunus dulcis | 1592.93 | 1592.93 | 99.77 | 0.0 | 100.0 | 157723 | NC\_034696.1 |
| Prunus dulcis chloroplast, complete genome | Prunus dulcis | 1592.93 | 1592.93 | 99.77 | 0.0 | 100.0 | 158085 | MT019559.1 |
| Prunus dulcis matK (matK) gene, complete cds; chloroplast gene for chloroplast product | Prunus dulcis | 1592.93 | 1592.93 | 99.77 | 0.0 | 100.0 | 1521 | AF288115.1 |
| Prunus triloba voucher Amygdalus triloba chloroplast, complete genome >gi|1817956713|gb|MK790138.1| Prunus triloba chloroplast, complete genome | Prunus triloba | 1587.39 | 1587.39 | 99.77 | 0.0 | 99.88 | 158455 | NC\_046742.1 |
| Prunus pedunculata chloroplast, complete genome | Prunus pedunculata | 1587.39 | 1587.39 | 99.77 | 0.0 | 99.88 | 157830 | OR343251.1 |
| Prunus japonica chloroplast, complete genome | Prunus japonica | 1587.39 | 1587.39 | 99.77 | 0.0 | 99.88 | 157927 | OP598111.1 |
| Prunus tomentosa chloroplast, complete genome | Prunus tomentosa | 1587.39 | 1587.39 | 99.77 | 0.0 | 99.88 | 157517 | MK911759.1 |
| Prunus japonica var. nakaii chloroplast, complete genome | Prunus japonica | 1587.39 | 1587.39 | 99.77 | 0.0 | 99.88 | 158078 | OM272947.1 |
| Prunus tomentosa isolate ZFI207 chloroplast, complete genome | Prunus tomentosa | 1587.39 | 1587.39 | 99.77 | 0.0 | 99.88 | 157082 | MT576930.1 |
| Prunus tomentosa isolate SMG211 chloroplast, complete genome | Prunus tomentosa | 1587.39 | 1587.39 | 99.77 | 0.0 | 99.88 | 157250 | MT576919.1 |
| Prunus tomentosa isolate LD501 chloroplast, complete genome | Prunus tomentosa | 1587.39 | 1587.39 | 99.77 | 0.0 | 99.88 | 157019 | MT576914.1 |
| Prunus tianshanica isolate DXG3 chloroplast, complete genome >gi|2292911896|gb|MT576892.1| Prunus tianshanica isolate DXG3 chloroplast, complete genome | Prunus tianshanica | 1587.39 | 1587.39 | 99.77 | 0.0 | 99.88 | 157577 | NC\_066419.1 |
| Prunus pedunculata chloroplast, complete genome | Prunus pedunculata | 1587.39 | 1587.39 | 99.77 | 0.0 | 99.88 | 157873 | MT247725.1 |
| Prunus triloba isolate R17 chloroplast, complete genome | Prunus triloba | 1587.39 | 1587.39 | 99.77 | 0.0 | 99.88 | 157820 | MW477438.1 |
| Prunus tomentosa isolate R20 chloroplast, complete genome | Prunus tomentosa | 1587.39 | 1587.39 | 99.77 | 0.0 | 99.88 | 157571 | MW477437.1 |
| Prunus triloba cultivar Multiplex chloroplast, complete genome | Prunus triloba | 1587.39 | 1587.39 | 99.77 | 0.0 | 99.88 | 157827 | MT937181.1 |
| Prunus triloba chloroplast, complete genome | Prunus triloba | 1587.39 | 1587.39 | 99.77 | 0.0 | 99.88 | 157816 | MH748555.1 |

specific

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| Description | Scientific Name | Max Score | Total Score | Query Cover | E value | Per. Ident | Acc. Len | Accession |
| Hypericum androsaemum isolate BPTPS187 maturase K (matK) gene, complete cds; chloroplast | Hypericum androsaemum | 377.837 | 377.837 | 99.19 | 5.95779e-107 | 75.28 | 1542 | ON684601.1 |
| Hypericum androsaemum isolate BPTPS150 maturase K (matK) gene, complete cds; chloroplast | Hypericum androsaemum | 377.837 | 377.837 | 99.19 | 5.95779e-107 | 75.28 | 1542 | ON684591.1 |
| Hypericum androsaemum maturase K (matK) gene, partial cds; chloroplast | Hypericum androsaemum | 377.837 | 377.837 | 99.19 | 5.95779e-107 | 75.28 | 1017 | HQ331618.1 |