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RcTric 1 -----MEQQQQGNGVVMGKRFIQQIQAQKMKELNNGFQSMWKSQKSLAVEAAVVTATSATGGAAGTACMGILAPDISSSLTPPPD--HALNPQAMASHKQQAALGGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
SmTric 1 -----GSLTEMAVAVERLQCLLAGRQWENQSAFVEMVMTAGCAAGGATGCGMGTADAAAGAPMGHQLPTNPDQAVVNLDFQNGGPPQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
AtTric1 1 --MVVGGGGGGDKRRSSGEMMAMASLFDNDQNIQDFVQKAEVITNFKWMSKQSTPVEAAVVTATSGELGGAIGMGLIGLSPMP-----Q-HGVDVPOAMASHKQQAALGGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
AtTric2 1 ---MGKDGEGDKKR---ETMAVMSLMKDOONPIQDFVQKAEVITNFKWMSKQSTPVEAAVVTATSGELGGAIGMGLIGLSPMP-----Q-HGVDVPOAMASHKQQAALGGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
BrTric1 1 ---MGKDGEGDKKRGGDETHAVMSLMKDOONPIQDFVQKAEVITNFKWMSKQSTPVEAAVVTATSGELGGAIGMGLIGLSPMP-----Q-HGVDVPOAMASHKQQAALGGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
BrTric2 1 ---MGVGGEGEGRRGEMMAISSLFDNDQNIQDFVQKAEVITNFKWMSKQSTPVEAAVVTATSGELGGAIGMGLIGLSPMP-----Q-7--DPOAMASHKQQAALGGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
BrTric3 1 -----MALIGGERRVTSGGDTHESSMKRISPTQYKIVSIVTFMTPGGVSGVAVLGGIIGTVMA-----PMHRRYPQQAALRRTOYA---RAYVLEGSVAIFAAGFIESMRRGIRG--KDDVTSRIVSASGGG
BrTric4 1 -----MGKDGDKVQETHAVMSLMKDOONPIQDFVQKAEVITNFKWMSKQSTPVEAAVVTATSGELGGAIGMGLIGLSPMP-----Q-HGVDVPOAMASHKQQAALGGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
CruTric1 1 --MGVGGGGGGDQRRGGGEMMAMASLFDNDQNIQDFVQKAEVITNFKWMSKQSTPVEAAVVTATSGELGGAIGMGLIGLSPMP-----Q-HGVDVPOAMASHKQQAALGGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
CruTric2 1 -----MAPLFAVAVPLGVESSFDPEKQKHWADWDQVAVVAVLGLQSSVGGVAGIEMGFSISAID--PTNGTAPRPNALDSSMMKGGPQWQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
CreTric 1 -----MEQGGQGVVVKLPP---HONVPEIQIQAQKMKELNNGFQSMWKSQKSLAVEAAVVTATSATGGAAGTACMGILAPDISSSLTPPPD--HALNPQAMASHKQQAALGGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
EgTric1 1 -----MEREKGVMTVSKLAP---QONPEIQIQAQKMKELNNGFQSMWKSQKSLAVEAAVVTATSATGGAAGTACMGILAPDISSSLTPPPD--HALNPQAMASHKQQAALGGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
EgTric2 1 -----MEQKGRGIVVSKLPP---PNNPEIQIQAQKMKELNNGFQSMWKSQKSLAVEAAVVTATSATGGAAGTACMGILAPDISSSLTPPPD--HALNPQAMASHKQQAALGGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
EgTric3 1 -----MEQKGRGIVVSKLPP---PNNPEIQIQAQKMKELNNGFQSMWKSQKSLAVEAAVVTATSATGGAAGTACMGILAPDISSSLTPPPD--HALNPQAMASHKQQAALGGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
GmTric1 1 -----MEQAKQGIHVTS---OSQNPPEIQIQAQKMKELNNGFQSMWKSQKSLAVEAAVVTATSATGGAAGTACMGILAPDISSSLTPPPD--HALNPQAMASHKQQAALGGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
GmTric2 1 -----MEQAKQGIHVTS---OSQNPPEIQIQAQKMKELNNGFQSMWKSQKSLAVEAAVVTATSATGGAAGTACMGILAPDISSSLTPPPD--HALNPQAMASHKQQAALGGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
ZmTric1 1 MADGKGRNSAVALKTSP--SVPMARGLRGGSDPEEWSGKRVKAIAGFRAMNAKQPTHEAAVVTATSGVAGVGGAGLADGGSPFIQPP--SNANSDAMASFKQAQALAGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
ZmTric2 1 --MERAGQRRPLTVMSSSSQVAARMGMTNPLAEWDRVRSLEAGFRAMNAKQPTHEAAVVTATSGVAGVGGAGLADGGSPFIQPP--PGTDKRVVLSFKQAQALAGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
ZmTric3 1 MADGKGRNSAVALKTSPSSSVAAGLGRVSDPEEWSGKRVKAIAGFRAMNAKQPTHEAAVVTATSGVAGVGGAGLADGGSPFIQPP--FNANRPMASFKQAQALAGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
VvTric1 1 -----MDEGKQGVMAAQNPIQAQKMKELNNGFQSMWKSQKSLAVEAAVVTATSATGGAAGTACMGILAPDISSSLTPPPD--HALNPQAMASHKQQAALAGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
Ostric1 1 -----MEGMGQQRRLVVMAS---AQAAARGGANLAELELDRVRLTLAGLFRAMNAKQPTHEAAVVTATSGVAGVGGAGLADGGSPFIQPP--PGADNPMASFKQAQALAGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
Ostric2 1 -----MEGKQGVMAVTKLPP---QONPEIQIQAQKMKELNNGFQSMWKSQKSLAVEAAVVTATSATGGAAGTACMGILAPDISSSLTPPPD--HALNPQAMASHKQQAALAGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
StTric 1 -----MEGKQGVMAVTKLPP---QONPEIQIQAQKMKELNNGFQSMWKSQKSLAVEAAVVTATSATGGAAGTACMGILAPDISSSLTPPPD--HALNPQAMASHKQQAALAGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
PpTric1 1 -----MQAATNKADGNTQMIQLKVKLSDDLVWVKKQPAAEVAVVVTAGVAGVGGAGLADGGSPFIQPP--FOLNPMASFKQAQALAGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
PpTric2 1 -----MEQKQGVVTKLPP---NDDPKVLFQKVKELDGGKSMWKSQKSLAVEAAVVTATSGVAGVGGAGLADGGSPFIQPP--HALNPQAMASHKQQAALAGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
PpTric3 1 -----MEQKQGVVTKLPP---NDDPKVLFQKVKELDGGKSMWKSQKSLAVEAAVVTATSGVAGVGGAGLADGGSPFIQPP--HALNPQAMASHKQQAALAGGLVQARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG
VcTric 1 -----MALPFAVAVPLGVEVKSDELGKRYQDWLAGQDVAVQVLLGLQSGVAVGGVGVVMPGFSISFD--PNNQ-GAAPSNPALESVVKAGPQSGARNFVMTGYNAGISSVKKRLRG--KEDVQSSMVAAFGSG

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RcTric 145 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
SmTric 133 AGLFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RSPRAHIV-----
AtTric1 149 FAYSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
AtTric2 144 FAYSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
BrTric1 148 FAYSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
BrTric2 144 FAYSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
BrTric3 121 LAYSFLAV--KGLTGGFAHAFPSAYVVAASG---TATIKSRNAODAFVTEKNAISKLGLEEKYKKNFKKGLLDDTLPLLLTDS--VLOQVIPPGRRLIILDHIQ--RFYKLSLYSMDQIYET-----
BrTric4 21 LAYSFLAV--KGLKVPFAHAFPSAYVVAASG---TATIKSRNAODAFVTEKNAISKLGLEEKYKKNFKKGLLDDTLPLLLTDS--VLOQVIPPGRRLIILDHIQ--RFYKLSLYSMDQIYET-----
CruTric1 143 FAYSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
CruTric2 150 FAYSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
CreTric 130 VAYSFLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RNPVSVLRFALPLPLPPPPPPMAAAAGASGR
EgTric1 145 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
EgTric2 144 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
EgTric3 9 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
GmTric1 144 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
GmTric2 144 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
GmTric3 158 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
ZmTric1 158 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
ZmTric2 158 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
ZmTric3 159 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
VvTric1 56 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
VvTric2 140 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
Ostric1 60 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
Ostric2 154 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
MTric 147 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
StTric 147 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
PpTric1 78 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
PpTric2 137 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
PpTric3 147 NMFSLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RDPELKERRRGHG-----
VcTric 128 VAYSFLVSGMGGPNQANAVTSGLFFALVGGGLVQGGKQKFSOPPAEDLDFVSRTRSMLSLGLQNKYKKNFKKGLLDDTLPLLLTDS--ALRDVIPPGRRLIILDHIQ--RNPVSVLRFALPLPLPPPPPPMAAAAGASGR

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Figure S1. Protein sequence alignment. Multiple sequence alignment (Clustal Omega) of all Tric orthologs from 17 plant species and algae (red, green and brown), with plant species chosen as representatives of each evolutionary clade from *Physcomitrella patens* (Pp) to *Eucalyptus grandis*, (Eg). Abbreviations: Br = *Brassica rapa*, Cru = *Capsella rubella*, Cre = *Chlamydomonas reinhardtii*, Cs = *Cucumis sativus*, Dm = *Drosophila melanogaster*, Eg = *Eucalyptus grandis*, Gm = *Glycine max*, Mt = *Medicago truncatula*, Os = *Oryza sativa*, Pp = *Physcomitrella patens*, Pt = *Populus Trichocarpa*, Rc = *Ricinus communis*, Sc = *Saccharomyces cerevisiae*, Sm = *Selaginella moellendorffii*, St = *Solanum tuberosum*, Vv = *Vitis Vinifera*, Vc = *Volvox carteri*, Zm = *Zea mays*.