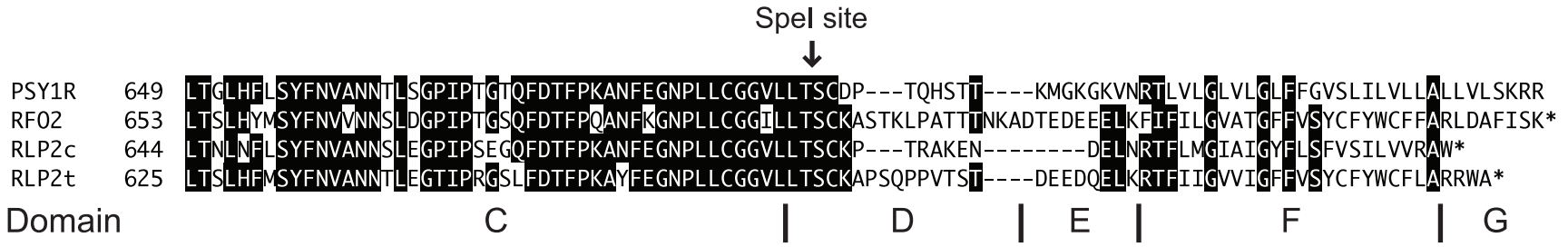
SpeI site
PSY1R 649 LTGLHFLSYFNVANNTLSGPIPTGTQFDTFPKANFEGNPLLCGGVLLTSCDP---TQHSTT----KMGKGKVNRTLVLGLVLGLFFGVSLILVLLALLVLSKRR
RFO2 653 LTSLHYMSYFNVVNNSLDGPIPTGSQFDTFPQANFKGNPLLCGGILLTSCKASTKLPATTTNKADTEDEEELKFIFILGVATGFFVSYCFYWCFFARLDAFISK*
RLP2c 644 LTNLNFLSYFNVANNSLEGPIPSEGQFDTFPKANFEGNPLLCGGVLLTSCKP---TRAKEN--------DELNRTFLMGIAIGYFLSFVSILVVRAW*
RLP2t 625 LTSLHFMSYFNVANNTLEGTIPRGSLFDTFPKAYFEGNPLLCGGVLLTSCKAPSQPPVTST----DEEDQELKRTFIIGVVIGFFVSYCFYWCFLARRWA*
Domain C D E F G