```
agrin_triad_egf1  C...RTFRCFQNGQC...LNRTFGYYTCQCPLGYTGTYC
agrin_triad_egf2  C...SPNPCQNGGGC.SPLSG.NRYFQCACPTDYAGLLC
agrin_triad_egf3  C...ASNPCPANSTC.THEAGKSDSWKCLCPLGKIGQRC
agrin_triad_egf4  CQR...KPCQNGAQC..SLGVQG..YLCNCPQYYTGENC
agrin_strpu_egf1  C...DSSPCQHGGTC.QNDEI.APGFRCICPLGKGGPVC
agrin_strpu_egf2  C.QSKEMPCFNNGLC.EALNAES..YRCICQGDFFGTLC
agrin_strpu_egf3  C...EGHMCHEESTC.VALPE.GG.YRCDCPDGRMGDMC
agrin_cioin_egf1  CI...PNPCQHSANCFITMQANIFTNKCECKDNYEGETC
agrin_cioin_egf2  C...SPNPCQGGAKC.IEMPG.EEEFTCKCPPGRSGSLC
agrin_cioin_egf3  CYR...NPCDNGGVC..HPRGAE..YMCVCLPYYTGDNC
agrin_rat_egf1    C...DSQPCLHGGTC.QDQDS.GKGFTCSCTAGRGGSVC
agrin_rat_egf2    C...LPNPCHGGALC.QALEA.GM.FLCQCPPGRFGPTC
agrin_rat_egf3    C...QPNPCHGAAPC.RVLSS.GG.AKCECPLGRSGTFC
agrin_rat_egf4    CTQALGNPCLNGGSC..VPREAT..YECLCPGGFSGLHC
```