

AtS8H	Sequence	102030405060
	Sequence	MGINFEDQTT LFNFVVR <del>EG</del> N GVKGMID <del>S</del> GL SSVPRPFVQP LSE <del>R</del> IP <del>T</del> QKA LTC <del>E</del> ATQ <del>P</del> I <del>D</del>
Potri.005G097900.1	Sequence	MAPSFLD <del>G</del> NS MFNFVVRD <del>G</del> N GVKGMVD <del>F</del> GL SEVPEQYVQP PEERID <del>K</del> LNA TAH <del>D</del> N-PPID
AtS8H	Sequence	708090100110120
	Sequence	LSNLDG <del>P</del> QHK EVAKQIV <del>E</del> AA ETLGFFQV <del>V</del> N HGVSV <del>E</del> LLEL LKSSA <del>H</del> EFFA QAP <del>E</del> EKSMYL
Potri.005G097900.1	Sequence	LSKLDG <del>P</del> DHD QIV <del>E</del> EIATAA ETLGFFQV <del>M</del> N HGV <del>P</del> V <del>E</del> LLES LKDAAN <del>N</del> FFG QPPE <del>K</del> KAVYR
AtS8H	Sequence	130140150160170180
	Sequence	KEVSPSKLVK YGTSFVPD <del>K</del> E KAI <del>E</del> WKDYVS MLYTND <del>S</del> EAL QHWPQPC <del>R</del> EV AL <del>E</del> FLNSSME
Potri.005G097900.1	Sequence	KGVSPSPSVR YGTSFVPD <del>K</del> E KAL <del>E</del> WKDYIS MRYTTDA <del>E</del> AL EYWPQEC <del>K</del> DV AL <del>E</del> YLRTSIK
AtS8H	Sequence	190200210220230240
	Sequence	MVKNVVNILM ENVGV <del>T</del> LEEE KMNGLMGTKM VNMNY <del>Y</del> PTCP SPELTVGVGR HSDMGMLT <del>V</del> L
Potri.005G097900.1	Sequence	MVRKVLEILI GKLG <del>V</del> TLDDS KIDGLIGLKL VNMN <del>F</del> YPTCP NPELTVGVGR HSDMGILT <del>V</del> L
AtS8H	Sequence	250260270280290300
	Sequence	LQDGIGGLYV KLD----- NG <del>E</del> WAEIPPV HGALVINIGD TLQILSNGKY KSAEHRVRTT
Potri.005G097900.1	Sequence	LQDDIGGLYV KMEEEMDGKR KGDWLEIPPV PGALVINVGD TLQILSNGRY KSAEHRVRTT
AtS8H	Sequence	310320330340350360
	Sequence	NIGSRVSVPI FTAPNPSQKV GPLPEVVKRD GVARYKEFLF QDYMN <del>N</del> FFGQ PHDGKKSLDF
Potri.005G097900.1	Sequence	RTKSRVSIPI FTIPKPT <del>E</del> RI GPLQVVERD GVARYREFIF EEYMN <del>N</del> FFSN AHDGKKSLDF
AtS8H	Sequence	370380390400410420
	Sequence	ARAE---. . . . .
Potri.005G097900.1	Sequence	AKNIWPA. . . . .