

MMETSP0253-20130528|12949_1 PSKLMPPKGVVKGKCAAKVARGRGAKALVLNGRRERTAGGLKADML MRNRSGKVSKRASA
DVNP2_sequenced MRNRSGKVSKRASA

MMETSP0253-20130528|12949_1 AGRRNFAK IEPWLQAVMTARECLRVTGFVAINGRTLQ GKALYVKSRAIYGSGV KGTEADP
DVNP2_sequenced AGRRNFA T IEPWLQAVMTARECLRVTGFVAINGRTLQ GKALYVKSRAIYGSGF KGTEADP

MMETSP0253-20130528|12949_1 VSSSAAASAG GDT M SSLAKRGCS D GGLILNW RV V V RGWSSGHICCC
DVNP2_sequenced VSSSAAASAG AHF V ELGEAGL.....FWRP H L E L AR.....