



Log<sub>2</sub> fold changes of probes  
[Nucleotide excision repair; n = 10]

	<u>LL<sub>3</sub></u> DD <sub>3</sub>	<u>Diel<sub>3</sub></u> LL <sub>3</sub>	<u>DD<sub>3</sub></u> Diel <sub>3</sub>	<u>LL<sub>6</sub></u> DD <sub>6</sub>	<u>Diel<sub>6</sub></u> LL <sub>6</sub>	<u>DD<sub>6</sub></u> Diel <sub>6</sub>	
K03723 mfd; transcription–repair coupling factor (superfamily II helicase) [EC:5.6.2.4]							SAR116_0227 transcription–repair coupling factor
K03701 uvrA; excinuclease ABC subunit A							SAR116_0401 excinuclease ABC
K03702 uvrB; excinuclease ABC subunit B							SAR116_2210 excinuclease ABC
K03703 uvrC; excinuclease ABC subunit C							SAR116_2209 Nuclease subunit of the excinuclease complex
K03657 uvrD; ATP–dependent DNA helicase UvrD/PcrA [EC:5.6.2.4]							SAR116_2321 DNA helicase II
K02335 polA; DNA polymerase I [EC:2.7.7.7]							SAR116_1268 DNA polymerase I
K02335 polA; DNA polymerase I [EC:2.7.7.7]							SAR116_0390 putative phage exonuclease
K02335 polA; DNA polymerase I [EC:2.7.7.7]							SAR116_0391 DNA polymerase
K02335 polA; DNA polymerase I [EC:2.7.7.7]							SAR116_1520 DNA polymerase I – 3'–5' exonuclease and polymerase domains
K01972 E6.5.1.2; DNA ligase (NAD+) [EC:6.5.1.2]							SAR116_2323 DNA ligase (NAD+)