



logo

midas g 534, midas s 5291, Hjor.155	CDDGGT	171
Bryobacter, midas s 6438, Kalu.261	LDDGA.	171
Bryobacter, midas s 3298, Fred.662 (split)	LDDGK.	191
Paludibaculum, midas s 675, Skiv.396 (split)	LDDGR.	189
Komagataeibacter xylinus, QUERY	LDDGR.	191
Rhodoferax, midas s 901, Aved.394	LDDGR.	207
Rhodoferax, midas s 901, EsbW.485 (merged)	LDDGR.	207
Dechloromonas, midas s 173, EsbW.050	LDDGR.	312
Dechloromonas, midas s 173, Hade.364	LDDGR.	312
Dechloromonas, midas s 173, Skiv.078 sub	LDDGR.	312
Nitrotoga, midas s 181, OdNE.187	LDDGR.	300
Nitrotoga, midas s 181, Skiv.176 (merged)	LDDGR.	300
Simplicispira, Simplicispira limi, EsbW.453	LDDGK.	276
Rhodoferax, midas s 4982, Bjer.458 A	LDDGR.	310
Rhodoferax, midas s 4982, Bjer.458 B	LDDGR.	19
consensus	*!!!*	

 non-conserved  
 ≥ 50% conserved