

## A

Construct Name	Primer Pair	Restriction Enzyme	Template
1:WT	T6WT Fwd x T6WT Rev	(BamHI/HindIII)	MGAS10394 genomic DNA
2:WT minus extension	T6noExt Fwd x T6noExt Rev	(BamHI/HindIII)	1:WT T6
3:N and M domains	T6Ndom Fwd x T6Mdom Rev	(KasI/XhoI)	1:WT T6
4:M	T6Mdom Fwd x T6Mdom Rev	(KasI/XhoI)	1:WT T6
5:M and C domains + ext	T6Mdom Fwd x T6Cdom Rev	(KasI/XhoI)	1:WT T6
6:M and C domains - ext	T6Mdom Fwd x T6Cdom Rev	(KasI/XhoI)	5:WT minus extension
7:M1	M1_0128 Fwd x M1_0128 Rev	(BamHI/EcoRI)	SF370 genomic DNA
8:M2	M2_0109 Fwd x M2_0109 Rev	(BamHI/HindIII)	MGAS10270 genomic DNA
9:M4	M4_0116 Fwd x M4_0116 Rev	(BamHI/EcoRI)	MGAS10750 genomic DNA
10:M28	M28_0109 Fwd x M28_0109 Rev	(BamHI/EcoRI)	MGAS6180 genomic DNA
11:M49	M49_FctA Fwd x M49_FctA Rev	(BamHI/EcoRI)	94/229* genomic DNA

## B

Primer Name	Primer Sequence
T6WT Fwd	<u>cgggatcctt</u> atcaaaagatgatactgcacaac
T6WT Rev	cgaagcttttatccacctgtcgaaggaattcacc
T6Ndom Fwd	<u>gtaaaggcgcc</u> aaagatgatactgcacaactaaag
T6Mdom Fwd	<u>gtaaaggcgccc</u> atctattacaaagaaagtaacc
T6Mdom Rev	gtaaactcgcagttaataagtataaacaattttagaatcttc
T6Cdom Rev	gtaaactcgcagttatagcttggttagggatgtcag
T6noExt Fwd	Ⓣ-ggaggttcaggtacagtattattag
T6noExt Rev	Ⓣ-ccaatctgcccgaatctcataag
M1_0128 Fwd	<u>cgggatccc</u> gtacaacagttcacgg
M1_0128 Rev	cggaattcttattcaagacttttttatttg
M2_0109 Fwd	<u>cgggatccg</u> aggacaccagagtgcttctc
M2_0109 Rev	cgaagcttttaaccaccagattagggattgaagg
M4_0116 Fwd	<u>gcggatccg</u> aatcatcacataaaaccgatg
M4_0116 Rev	cggaattcttaaccaccagttgaggatgg
M28_0109 Fwd	<u>gcggtatccg</u> agacggcaggagtgccgaaaatg
M28_0109 Rev	cggaattcttaaacaccagttggaacttgagtg
M49_FctA Fwd	<u>ctagggatccg</u> agacagcaggagtgattg
M49_FctA Rev	cggaattcttaaacaccagttggaacttgagtg

**TABLE S1**

(A) Construct details of the clones used in this work. (B) Primer sequences are shown with the gene specific sequence as Black text and the non-gene specific sequence as Grey text with the restriction site underlined. Ⓣ- 5' phosphate group. \* Proft *et al.* (2000) (1).

1. **Proft T, Moffatt SL, Weller KD, Paterson A, Martin D, Fraser JD. 2000.** The streptococcal superantigen SMEZ exhibits wide allelic variation, mosaic structure, and significant antigenic variation. *J Exp Med* **191**:1765-1776.