

Supplementary Figure S1: Clustal alignment of CSP amino acid haplotypes.

Deduced protein sequence of sequenced region including Th2R and Th3R epitopes, showing 29 different sequences observed among 81 ADI samples analysed, compared to 3D7 (vaccine antigen) sequence. Majority residues are white, minority substitutions are black.

3D7CSP	NANANS AVKNNNNEEPSDKH I KEYL N KIQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKD E LDYANDIEKKICKMEKCSV 0001 NANANNAE K NNNNEEPSDQHIEKYLKII IQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLDYANDIEKKICKMEKCSV 0002 NANANNAVKNNNNEEPSDQHIEKYLKII IQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLDYANDIEKKICKMEKCSV 0003 NANANNAVKNNNNEEPSDQHIEKYLQKIQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLNYENDIEKKICKMEKCSV 0004 NANANNAVKNNNNEEPSDQHIEQYLNRIQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLNYENDIEKKICKMEKCSV 0005 NAKANNAVKNNNNEEPSDKHIEQYLKTIQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLDYENDIEKKICKMEKCSV 0006 NANANNAVKNNNNEEPSDKHIEQYLKTIQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLDYENDIEKKICKMEKCSV 0007 NANANNAVKNNNNEEPSDQHIEQYLKTIQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLDYENDIEKKICKMEKCSV 0008 NANANNAVKNNNNEEPSDKHIEEYLKKIKNSISTEWSPCSVT CGNGI QVRIKPGSANKPKD E LN YENDIEKKICKMEKCSV 0009 NANANNAVKNNNNEEPSDKHIEQYLKKIKNSISTEWSPCSVT CGNGI QVRIKPGSANKPKD E LDYENDIEKKICKMEKCSV 0010 NANANNAVKNNNNEEPSDKHIEQYLKKIQNSLSTEWSPCSVT CGNGI QVRIKPGSA G SKD E LDYENDIEKKICKMEKCSV 0011 NANANNAVKNNNNEEPSDKHIEKYLKEIQNSLSTEWSPCSVT CGNGI QVRIKPGSA G KPK N ELDYENDIEKKICKMEKCSV 0012 NANANNAVKNNNNEEPSDQHIEKYLKKIQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLDYANDIEKKICKMEKCSV 0014 NANANNAVKNNNNEEPSDQHIEKYLKKIQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLDYENDIEKKICKMEKCSV 0015 NANANNAVKNNNNEEPSDKH I KEYLKII IQNSLSTEWSPCSVT CGNGI QVRIKPGSA G KPK E LN YENDIEKKICKMEKCSV 0016 NANANNAVKNNNNEEPSDKHIEQYLKTIKNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLDYENDIEKKICKMEKCSV 0017 NANANNAVKNNNNEEPSDQHIEQYLKTIKNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLDYANDIEKKICKMEKCSV 0018 NANANNAVKNNNNEEPSDQHIEKYLKII IQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLDYANDIEKKICKMEKCSV 0019 NANANNAVKNNNNEEPSDQHIEKYLQKIQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLDYENDIEKKICKMEKCSV 0020 NANANNAVKNNNNEEPSDQHIEKYLNKIKNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLDYANDIEKKICKMEKCSV 0021 NANANNAVKNNNNEEPSDQHIEKYLKII IQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLDYANDIEKKICKMEKCSV 0022 NANANNAVKNNNNEEPSDKHIEQYLKKIQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKD E LDYENDIEKKICKMEKCSV 0023 NANANNAVKNNNNEEPSDKHIEQYLKKIKNSISTEWSPCSVT CGNGI QVRIKPGSANKPKD E LDYENDIEKKICKMEKCSV 0024 NANANNAVKNNNNEEPSDKHIEQYLKKIQNSLSTEWSPCSVT CGNGI QVRIKPGSA G SKD E LDYENDIEKKICKMEKCSV 0025 NANANNAVKNNNNEEPSDKHIEEYLKKIKNSISTEWSPCSVT CGNGI QVRIKPGSANKPKD E LN YENDIEKKICKMEKCSV 0026 NANANNAVKNNNNEEPSDQHIEKYLQKIQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLNYENDIEKKICKMEKCSV 0027 NANANNAVKNNNNEEPSDQHIEQYLKRIQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLDYANDIEKKICKMEKCSV 0028 NAKANNAVKNNNNEEPSDKHIEQYLKTIQNSLSTEWSPCSVT CGNGI QVRIKPGSANKPKDQLDYENDIEKKICKMEKCSV 0029 NANANNAVKNNNNEEPSDKH I KEYLKII IQNSLSTEWSPCSVT CGNGI QVRIKPGSA G KPK F ELNYENDIEKKICKMEKCSV 0030 NANANNAVKNNNNEEPSDKHIEKYLKEIQNSLSTEWSPCSVT CGNGI QVRIKPGSA G KPK N ELDYENDIEKKICKMEKCSV
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Th2R

Th3R