

SARS-COV-2 VARIANTS IN RISK ASSESSMENT: VARIANTS DETECTED IN THE UK

POSITION AS OF 07 July 2021 PHE

Last horizon scan	Signals currently under monitoring and investigation	Variants under investigation	Variants of concern
International Epidemiology Scan: 06-07-2021	Epsilon B.1.427/B.1.429 (California)	Zeta - VUI-21JAN-01 P.2 (Brazil)	Alpha - VOC-20DEC-01 B.1.1.7 (UK)
Phylogenetic Cluster detection: 05-07-2021	Iota B.1.526 (New York)	Eta - VUI-21FEB-03 B.1.525	Beta - VOC-20DEC-02 B.1.351 (South Africa)
Imported Sample Mutation Scan: 05-07-2021	B.1.1.7 with S494P (UK)	VUI-21FEB-04 B.1.1.318 (England)	Gamma - VOC-21JAN-02 P.1 (Japan ex Brazil)
Signals in Monitoring Scan: 06-07-2021	A.27 (France)	Theta - VUI-21MAR-02 P.3 (Philippines)	Delta - VOC-21APR-02 B.1.617.2 (India), AY.1, and AY.2
International Mutation Scan: 06-07-2021	B.1.1.7 with Q677H (UK)	Kappa - VUI-21APR-01 B.1.617.1 (India)	
Domestic Mutation Scan: 02-07-2021	B.1.620 (travellers to the UK)	VUI-21APR-03 B.1.617.3 (India)	
Reinfections Scan: 25-06-2021	B1.214.2 (Belgium)	VUI-21MAY-01 AV.1 (UK)	
Vaccinees Scan: 02-07-2021	R.1 Associated with multiple locations	VUI-21MAY-02 C.36.3 (Egypt)	
	B.1.621 (Colombia)	Lambda – VUI-21JUN-01 C.37 (Peru)	
	AT.1 (Russia)		
Signals investigated at last scan: 0	Delta like variant with E484A (India, UK, Japan)		
New genomes in last 24 hr period: 5119	Lineage A with R346K, T478R and E484K (Tanzania/Angola)		
Note: Signals of individual mutations and new clusters excluded until defined variant identified	B.1.629 (Guinea/Angola)		
	B.1.619 (S. Korea)		

Variants provisionally extinct (not detected for 12 weeks in the UK)	International signals under monitoring	International variants under investigation	International variants of concern
VUI-21MAR-01 B.1.324.1 with E484K (UK) VUI-21FEB-01, A.23.1 with E484K, (UK) focus Liverpool	P.1 + N501T and E484Q (Brazil)		VOC-21FEB-02 B.1.1.7 with E484K (UK) focus Bristol/SW.
	Variant Associated with North America: USA 214insQAS Samples		

MUTATION PROFILES: VARIANTS OF CONCERN AND VARIANTS UNDER INVESTIGATION

VARIANT	MUTATION PROFILE (SUBSTITUTIONS AND INSERTION/DELETIONS)
Alpha VOC-20DEC-01	S: Δ69-70, Δ144, N501Y, A570D, P681H, T716I, S982A, D1118H; ORF1ab: T1001I, A1708D, I2230T, Δ3675-3677; ORF8: Q27*, R52I, Y73C; N: D3L
Beta VOC-20DEC-02	S: D80A, D215G, Δ242-244, K417N, E484K, N501Y, A701V; ORF1ab: T265I, K1655N, K3353R, Δ3675-3677; ORF3a: Q57H, S171L; E: P71L, T205I; N:T205I
Gamma VOC-21JAN-02	S: T20N, P26S, K417T, E484K N501Y, T1027I; ORF1ab: S1188L, K1795Q, Δ3675-3677; ORF8: E92K; N: P80R
VUI-21FEB-01	S: R102I, F157L, V367F, E484K, Q613H, P681R; ORF1ab: L1559F, M3655I, L3667F, M3752I; ORF8: L84S, E92K; N: S202N
Eta VUI-21FEB-03	S: Q52R, Δ69-70, Δ144, E484K, Q677H, F888L; ORF1ab: Δ3675-3677, P4715F; E: L21F; M: I82T; ORF6: Δ2; N: Δ2, D3Y
VUI-21FEB-04	S: T95I, Δ144, E484K, P681H, D796H; ORF1ab: E1196V, K2511N, T2936I, A3209V, T3284I, Δ3675-3677, V6672M; M: I82T; ORF8: Δ1-3, E106*; N: Δ208, R209G
Zeta VUI-21JAN-01	S: E484K; ORF1ab: L3468V; N: A119S, M234I
VUI-21MAR-01	B.1.324.1 + S: E484K; ORF1ab: G894S, G5530C; ORF8: Δ27922-27956
Theta VUI-21MAR-02	S: Δ141-143, E484K, N501Y, P681H, E1092K, H1101Y, V1176F; ORF1ab: L71F, D112E, A368V, L438P, D736G; ORF8: K2Q
Kappa VUI-21APR-01	S: E154K, L452R, E484Q, P681R; ORF1ab: T1567I, T3646A, M5753I, K6711R; ORF7a: V82A; N: R203M
Delta VOC-21APR-02	B.1.617.2: S: T19R, G142D, Δ156-157, R158G, L452R, T478K, P681R, D950N; ORF3a: S26L; M: I82T, ORF7a: V82A, T120I N: D63G, R203M, D377Y AY.1: S:T19R, G142D, Δ156-157, W258L, K417N , L452R, T478K, D614G, P681R, D950N; ORF 3a:S26L; M:I82T; ORF 7a:V82A, T120I; ORF 7b:T40I ; N:D63G, G215C, R203M, D377Y

Where INDEL are defined in 'nt' the INDEL is non-codon aligned.

Note: These are not lineage defining mutations, these are mutations found in the VOC/VUIs.

Lineage definitions can be found [here](#)

MUTATION PROFILES: VARIANTS OF CONCERN AND VARIANTS UNDER INVESTIGATION

VARIANT	MUTATION PROFILE (SUBSTITUTIONS AND INSERTION/DELETIONS)
VUI-21APR-03	S: T19R, Δ156-157, R158G, L452R, E484Q, P681R, D950N, ORF1ab: A1526V, T1830I, A194S, A117V, ORF7a: V82A, ORF8: T26I, N: P67S, R203M, D377Y
VUI-21MAY-01	S: D80G, T95I, G142D, Δ144, N439K, E484K, D614G, P681H, I1130V, D1139H , Orf1ab: A119V, G339S, H342Y, G519S, A591V, H1160Y, P1640L, N2405S, A3209V, Δ3675-3677, P4715L, S4826A, A6044V, M: A63T, H125Y , Orf3a: Δ26158-26161 , N: I157V, R203K, G204R
VUI-21MAY-02	S: S12F, Δ69-70, W152R, R346S, L452R, Q677H, A899S ; orf1ab: D105Y, S118L, D217N, E102K, A859V, S3687L, L3691S, T1246I, D1639N, P2287S, D3222N, G3278S, T4090I, P4715L; N: R203K, G204R, G212V; M: I82T
Lambda VUI-21JUN-01	S: L452Q, F490S, T859N, Δ22298; ORF1ab: L438P, T492I, Δ11287, F1569V; N: P13L, G214C

Where INDEL are defined in 'nt' the INDEL is non-codon aligned.

MUTATION PROFILES: SIGNALS IN MONITORING

VARIANT	MUTATION PROFILE (SUBSTITUTIONS AND INSERTION/DELETIONS)
Epsilon B.1.427/B.1.429 (California)	S: S13I W152C L452R; N: T205I
Iota B.1.526 (New York)	S: L5F T95I D253G S477N/E484K A701V
B.1.1.7 with S494P (UK)	S: S494P
A.27 (France)	S: L18F L452R N501Y A653V H655Y D796Y G1219V
B.1.1.7 with Q677H (UK)	S: Q677H
B.1.620 (travellers to the UK)	S: P26S V126A S477N E484K P681H T1027I D1118H
R1 (Multiple Locations)	S: W152L E484K D614G G769V
B1.214.2 (Belgium)	S: Q414K N450K D614G T716I Ins: 22204 – ACAGATCGA
AT.1 (Russia)	S: D215G, E484K, E780K, H245P, P9L, Δ136-144, N679K/insGIAL
B.1.621 (Spain)	S: T95I, Y144T/144insS/Y145N*, R346K, E484K, N501Y, D614G, P681H, D950N
VOC-21FEB-02	VOC-20DEC-01 + S: E484K; N: A173V A398T
Lineage A +R346K, T478R, E484K (Angola/Tanzania)	S: D80Y, Δ144, Δ210, Δ246-248, L249M, D215G, R346K, T478R, E484K, H655Y, P681H, Q957H
Delta like variant with E484A (India, UK, Japan)	S: T76I, Δ144, D253N, L452R, T478K, E484A, T572N, D614G, P681R, D796H, T859N, S939F
B.1.629 (Guinea/Angola)	S: G142D, K182R, L452R, T478K, D614G, P681H, D796Y, C1235F
B.1.619 (S. Korea)	I210T, N440K, E484K, D614G, D936N, S939F, T1027I