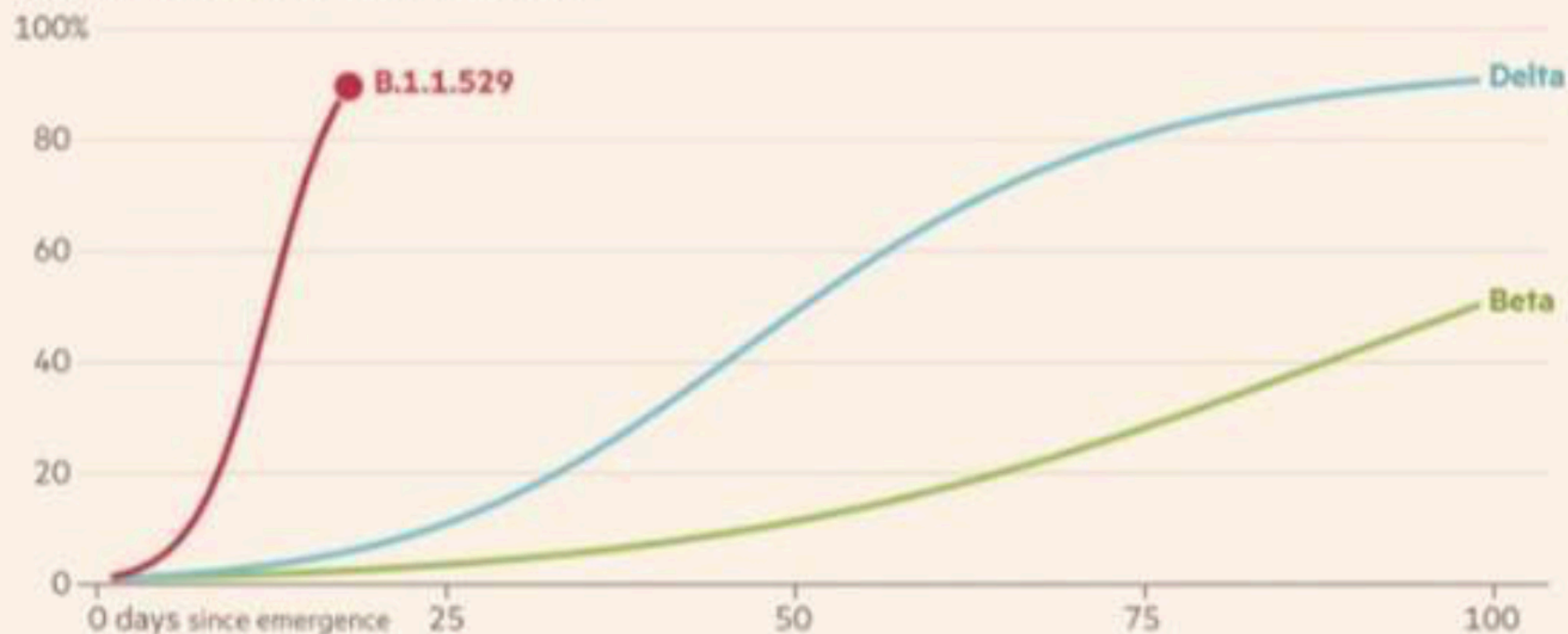


A new variant is spreading rapidly in South Africa, and appears to be out-competing other variants much faster than previous variants of concern did

Share of all sequenced cases\* in South Africa accounted for by each variant, by number of days since it passed 1%



\*Growth of B.1.1.529 is modelled from SGTf data rather than full genomic sequences

Source: FT analysis of data from GISAID and the South African National Health Laboratory Service

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