Table 3. De novo prediction of phylogroups, MLST types and serotypes of the sequenced UPEC isolates

Strain	<b>Accession Number</b>	Hospital	Phylogroups	MLST type	Serotype
NGE3	QEXN00000000	DCIMCH	F (D)*	ST-59	O1:H7
NGE4	QFAZ00000000	DCIMCH	Α	ST-4204	O6:H10
NGE5	RCIF00000000	DCIMCH	B1	ST-10987 (Ounknown ST)	O59:H20
NGE6	RCIE00000000	DCIMCH	E (B2)	ST-219	O138:H48
NGE7	QFRN00000000	DCIMCH	B2	ST-131	O46:H4 (O25:H4)
NGE9	QFRT00000000	DCIMCH	EC control fail (B2)	ST-219	O138:H48
NGE16	QFTM00000000	ISH	B2	NA (ST-131)	O25:H4
NGE22	QFXA00000000	ISH	Α	ST-4204	O6:H10
NGCE33	RBWA0000000	DCIMCH	C (A)	ST-410	O8:H9
NGCE94	RAZR00000000	DCIMCH	B1 (A)	ST-448 (ST-361)	O188:H19 (O9:H30)
NGCE100	RAZQ00000000	DCIMCH	A (B1)	ST-361 (ST-448)	H30:09 (Ounknown:H7)

<sup>\*</sup>Parentheses indicate originally published values that do not match with the reproduced values