# A new polymorphic probe on chromosome 22: NB17 (D22S181)

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*Source/Description*: NB17 is a 0.7 kb HindIII-EcoRI fragment isolated from two pooled chromosome 22 specific libraries (ATCC # 57733 and ATCC # 57714) and was subcloned into pUC9.

*Polymorphism*: BglII or TaqI digestion of genomic DNA and hybridization with the probe detects a two allele polymorphism: BglII (A1: 12.5 kb, A2: 4.0 kb) with a constant band at 3.0 kb; TaqI (B1: 2.9 kb, B2: 2.2 kb). The polymorphisms detected with these two enzymes are almost in complete linkage disequilibrium.

*Frequency*: Estimated from 92 (BglII) and 95 (TaqI) unrelated Caucasians.

A1: 0.69 B1: 0.65
A2: 0.31 B2: 0.35

*Not Polymorphic For*: BglI, DraI, MspI and PstI.

*Chromosomal Localization*: Regional localization was established by hybridization to a panel of somatic cell hybrids: PgMe-25Nu, containing only human chromosome 22; PgMo-22 and 1CB-17ANu, respectively containing both products of the Philadelphia translocation at 22q11; A3EW2-3B, containing the t(11;22) from Ewing's sarcoma (at 22q12) and 1/22 AM27 containing a t(1;22) at 22q13 (1, 2). The probe was assigned to chromosome 22 between 22pter and 22q11.

*Mendelian Inheritance*: Mendelian inheritance has been demonstrated in extended pedigrees of Gilles de la Tourette syndrome families (n = 380).

*Probe Availability*: Available for collaboration.

*References*: 1) Goyns,M.H., Young,B.D., Geurts van Kessel,A., de Klein,A., Grosveld,G., Bartram,C.R. and Bootsma,D. (1984) *Leukemia Res.* **8**, 547–553. 2) Geurts van Kessel,A., Turc-Carel,C., de Klein,A., Grosveld,G., Lenoir,G. and Bootsma,D. (1985) *Mol. Cell Biol.* **5**, 427–429.

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# A new polymorphic probe on chromosome 22: NB35 (D22S182)

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*Source/Description*: NB35 is a 0.8 kb HindIII-EcoRI fragment isolated from two pooled chromosome 22 specific libraries (ATCC # 57733 and ATCC # 57714) and was subcloned into pUC9.

*Polymorphism*: BglII digestion of genomic DNA and hybridization with the probe detects a two allele polymorphism: 8.7 kb (A1) and 8.0 kb (A2). No constant bands were present.

*Frequency*: Estimated from 96 unrelated Caucasians.
A1: 0.76 A2: 0.24.

*Not Polymorphic For*: BglI, DraI, MspI, PstI and TaqI.

*Chromosomal Localization*: Regional localization was established by hybridization to a panel of somatic cell hybrids: PgMe-25Nu, containing only human chromosome 22; PgMo-22 and 1CB-17ANu, respectively containing both products of the Philadelphia translocation at 22q11; A3EW2–3B, containing the t(11;22) from Ewing's sarcoma (at 22q12) and 1/22 AM27 containing a t(1;22) at 22q13 (1, 2). The probe was assigned to chromosome 22 between 22pter and 22q11.

*Mendelian Inheritance*: Mendelian inheritance has been demonstrated in extended pedigrees of Gilles de la Tourette syndrome families (n = 380).

*Probe Availability*: Available for collaboration.

*References*: 1) Goyns,M.H., Young,B.D., Geurts van Kessel,A., de Klein,A., Grosveld,G., Bartram,C.R. and Bootsma,D. (1984) *Leukemia Res.* **8**, 547–553. 2) Geurts van Kessel,A., Turc-Carel,C., de Klein,A., Grosveld,G., Lenoir,G. and Bootsma,D. (1985) *Mol. Cell Genet.* **5**, 427–429.

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