3. The method as claimed in claim 1 or claim 2, further comprising the step of replacing at least one constant domain of the heavy and/or light chain of the donor immunoglobulin with a constant domain of a heavy and/or light chain derived from an immunoglobulin derived from the target species.

4. The method as claimed in any preceding claim wherein the target species is a mammalian target species.

5. The method as claimed in claim 4 wherein the mammalian species is a companion animal selected from the group consisting of a dog, a cat and a horse.

6. The method as claimed in claim 4 wherein the mammalian species is a human.

7. A neutralising antibody or an antigen binding fragment thereof which is capable of specifically binding to human nerve growth factor (NGF) wherein the antibody or antigen binding fragment comprises a light chain variable region comprising the amino acid sequence of SEQ ID NO:13 or an amino acid sequence which has an identity of at least 85% thereto and a heavy chain variable region comprising the amino acid sequence of SEQ ID NO:14 or an amino acid sequence which has an identity of at least 85% thereto, wherein the antibody or an antigen binding fragment thereof does not contain any amino acid in any position within the framework regions of the heavy and light chain variable domains which would be foreign at that position in one or more immunoglobulins derived from a human, wherein the light chain variable region comprises a CDR1 sequence consisting of RASEDIYNALA, a CDR2 sequence consisting of NTDTLHT, and a CDR3 sequence consisting of HYFHYPRT and wherein the heavy chain variable region comprises a CDR1 sequence consisting of NNNVN, a CDR2 sequence consisting of GVWAGGATDYNSALK, and a CDR3 sequence consisting of DGGYSSSTLYAMDA.

8. The antibody or antigen binding fragment thereof as claimed in claim 7 wherein the light chain comprises the amino acid sequence of SEQ ID NO:25, or an amino acid sequence which has an identity of at least 85% thereto.

9. The antibody or antigen binding fragment thereof as claimed in claim 7 or claim 8 wherein the heavy chain comprises the amino acid sequence of SEQ ID NO:24, or an amino acid sequence which has a sequence identity of at least 85% thereto.