CLAIMS

1. A method for determining the suitability of a granulocyte for treating cancer, the method comprising:
   a. comparing a measured expression level of one or more genes by the granulocyte, wherein the one or more genes are associated with suitability for treating cancer and are selected from: *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PPP3CB, ANXA1, PERM, PLEC, ACSL1, RAC1,* and *PSMB2,* with the expression level of the same one or more genes in a reference standard; and
   b. determining the suitability of the granulocyte for treating cancer based on the comparison.

2. A method for identifying whether or not a donor produces granulocytes suitable for treating cancer, the method comprising:
   a. comparing a measured expression level of one or more genes by a granulocyte comprised in a sample obtainable from the donor, wherein the one or more genes are associated with suitability for treating cancer and are selected from: *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PPP3CB, ANXA1, PERM, PLEC, ACSL1, RAC1,* and *PSMB2,* with the expression level of the same one or more genes in a reference standard; and
   b. identifying whether or not the donor produces granulocytes suitable for treating cancer based on the comparison.

3. The method according to claim 1 or 2, wherein step b. comprises:

determining that the granulocyte is suitable for treating cancer or identifying that the donor produces granulocytes suitable for treating cancer when:

   i. the measured expression level of one or more of *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1,* and *PSMB2* is increased when compared to the reference standard when the reference standard is from a granulocyte unsuitable for treating cancer; or
   ii. the measured expression level of *ANXA1* and/or *PPP3CB* is decreased when compared to the reference standard, when the reference standard is from a granulocyte unsuitable for treating cancer; or

iii. the measured expression level of one or more of *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1,* and *PSMB2* is increased or the same when compared to the reference standard, when the reference standard is from a granulocyte suitable for treating cancer; or

iv. the measured expression level of *ANXA1* and/or *PPP3CB* is decreased or the same when compared to the reference standard, when the reference standard is from a granulocyte suitable for treating cancer; or

determining that the granulocyte is unsuitable for treating cancer or identifying that the donor produces granulocytes that are unsuitable for treating cancer when:

v. the measured expression level of one or more of *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1,* and *PSMB2* is decreased or the same when compared to the reference standard, when the reference standard is from a granulocyte unsuitable for treating cancer; or

vi. the measured expression level of *ANXA1* and/or *PPP3CB* is increased or the same when compared to the reference standard, when the reference standard is from a granulocyte unsuitable for treating cancer; or

vii. the measured expression level of one or more of *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1,* and *PSMB2* is decreased when compared to reference standard, when the reference standard is from a granulocyte suitable for treating cancer; or

viii. the measured expression level of *ANXA1* and/or *PPP3CB* is increased when compared to the reference standard, when the reference standard is from a granulocyte suitable for treating cancer.

4. A method for selecting whether or not a subject is suitable for treatment with a granulocyte or a stem cell for treating cancer, the method comprising:

a. comparing a measured expression level of one or more genes by a granulocyte comprised in a sample obtainable from the subject, wherein the one or more genes are associated with suitability for treating cancer and are selected from: *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PPP3CB, ANXA1, PERM, PLEC, ACSL1,*

RAC1, and *PSMB2* with the expression level of the same one or more genes in a reference standard; and

b. identifying whether or not the subject is suitable for treatment with a granulocyte or a stem cell for treating cancer based on the comparison.

5. A method for determining a subject's risk for developing cancer, the method comprising:

a. comparing a measured expression level of one or more genes by a granulocyte comprised in a sample obtainable from the subject, wherein the one or more genes are associated with suitability for treating cancer and are selected from: *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PPP3CB, ANXA1, PERM, PLEC, ACSL1, RAC1,* and *PSMB2* with the expression level of the same one or more genes in a reference standard; and

b. determining the subject's risk for developing cancer based on the comparison.

6. The method according to claim 4 or 5, wherein step b. comprises:

identifying the subject as suitable for treatment with a granulocyte or a stem cell for treating cancer or determining that the subject is at risk of developing cancer when:

i. the measured expression level of one or more of *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMKATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1,* and *PSMB2* is decreased or the same when compared to the reference standard, when the reference standard is from a granulocyte unsuitable for treating cancer; or

ii. the measured expression level of *ANXA1* and/or *PPP3CB* is increased or the same when compared to the reference standard, when the reference standard is from a granulocyte unsuitable for treating cancer; or

iii. the measured expression level of one or more of *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1,* and *PSMB2* is decreased when compared to the reference standard, when the reference standard is from a granulocyte suitable for treating cancer; or

iv. the measured expression level of *ANXA1* and/or *PPP3CB* is increased when compared to the reference standard, when the reference standard is from a granulocyte suitable for treating cancer; or

identifying the subject as unsuitable for treatment with a granulocyte or a stem cell for treating cancer when or determining that the subject is not at risk of developing cancer when:

v. the measured expression level of one or more of *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1,* and *PSMB2* is increased when compared to the reference standard when the reference standard is from a granulocyte unsuitable for treating cancer; or

vi. the measured expression level of *ANXA1* and/or *PPP3CB* is decreased when compared to the reference standard, when the reference standard is from a granulocyte unsuitable for treating cancer; or

vii. the measured expression level of one or more of *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1,* and *PSMB2* is increased or the same when compared to the reference standard, when the reference standard is from a granulocyte suitable for treating cancer; or

viii. the measured expression level of *ANXA1* and/or *PPP3CB* is decreased or the same when compared to the reference standard, when the reference standard is from a granulocyte suitable for treating cancer.

7. The method according any one of the preceding claims, wherein the method further comprises measuring the expression of the one or more genes.

8. The method according to any one of the preceding claims, wherein the granulocyte is a neutrophil.

9. The method according to any one of the preceding claims, further comprising measuring the expression level of one or more gene selected from: *S100A9* and *S100A8*.

10. The method according to any one of the preceding claims, wherein the one or more genes associated with suitability for treating cancer are one or more genes associated with:

a. killing cancer cells and are selected from: *GM2A, CTSG, CAP37, CYBB, GZMK, ATM, PERM, ACSL1, ATG7, SYK, DOCK8, RAC1, and PSMB2*; and/or

b. locating and/or binding to cancer cells and are selected from: *ANXA1, ITGB1, COMP, SLC2A1* and *PLEC*; and/or

c. recruitment of immune mediators and are selected from: *BCAP31, TAPBP, IKBKB,* and *PPP3CB*.

11. The method according to any one of the preceding claims, wherein the expression level is measured by proteomic techniques.

12. The method according to any one of the preceding claims, wherein the expression level is measured by transcriptomic techniques.

13. The method according to any one of claims 1-3 or 7-12 further comprising selecting a granulocyte when the granulocyte has been determined to be suitable for treating cancer or selecting a granulocyte from a sample obtainable from a donor when the donor has been identified as a donor that produces granulocytes for treating cancer, preferably wherein the selecting comprises isolating the granulocyte.

14. An *in vitro* method for obtaining a granulocyte for treating cancer, said method comprising obtaining a granulocyte from a sample obtainable from a donor wherein said donor produces granulocytes comprising:

a. increased expression of one or more of *ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, CTSG, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1, GM2A, CAP37,* and *PSMB2* when compared to a reference standard, wherein the reference standard is from a granulocyte unsuitable for treating cancer; and/or

b. decreased expression of *ANXA1* and/or *PPP3CB* when compared to a reference standard, wherein the reference standard is from a granulocyte unsuitable for treating cancer.

15. An *in vitro* method for obtaining a stem cell for treating cancer, said method comprising obtaining a stem cell from a sample obtainable from a donor wherein said donor produces granulocytes comprising:

a. increased expression of one or more of *ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, CTSG, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1, GM2A, CAP37,* and *PSMB2* when compared to a reference standard, wherein the reference standard is from a granulocyte unsuitable for treating cancer; and/or

b. decreased expression of *ANXA1* and/or *PPP3CB* when compared to a reference standard, wherein the reference standard is from a granulocyte unsuitable for treating cancer.

16. A method for preparing a stem cell suitable for treating cancer, the method comprising:
    a. identifying a donor that produces granulocytes suitable for treating cancer according to the method of any one of claims 2-3 or 7-13; and
    b. obtaining the stem cell from a sample obtainable from said donor.

17. A method for producing a stem cell for treating cancer, the method comprising:
    a. providing a stem cell obtainable from a sample from a donor wherein said donor produces granulocytes comprising:
        i. increased expression of one or more of *ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, CTSG, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1, GM2A, CAP37,* and *PSMB2* when compared to a reference standard, wherein the reference standard is from a granulocyte unsuitable for treating cancer; and/or
        ii. decreased expression of *ANXA1* and/or *PPP3CB* when compared to a reference standard, wherein the reference standard is from a granulocyte unsuitable for treating cancer ; and
    b. differentiating the stem cell into a different stem cell (preferably a precursor cell); and
    c. optionally isolating the different stem cell (preferably the precursor cell).

18. The method according to any one of claims 15-17 wherein the sample comprises a somatic cell and obtaining the stem cell from the sample comprises reprograming the somatic cell into a stem cell.

19. A method for isolating a granulocyte or stem cell for treating cancer, the method comprising:
    a. contacting a sample of granulocytes or a sample of stem cells with a binding means, wherein the binding means binds to one or more polypeptides selected from GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PPP3CB, ANXA1, PERM, PLEC, ACSL1, RAC1, and PSMB2; and

b. isolating the granulocyte or stem cell, respectively, based on the presence or absence of binding between the binding means and the one or more polypeptides.

20. A method for producing an engineered granulocyte or stem cell for treating cancer, the method comprising:

a. providing a granulocyte or stem cell, respectively; and

b. engineering the granulocyte or stem cell, respectively, to:

i. increase expression of one or more genes selected from: *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1,* and *PSMB2*; and/or

ii. decrease expression of *ANXA1* and/or *PPP3CB*;

thereby producing the engineered granulocyte or stem cell, respectively, wherein the engineered granulocyte or stem cell, respectively, is suitable for treating cancer.

21. A stem cell for treating cancer obtainable by the method according to any one of claims 15-20.

22. A stem cell which is capable of differentiating into a granulocyte for treating cancer, wherein the granulocyte comprises:

a. increased expression of one or more of *ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, CTSG, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1, GM2A, CAP37,* and *PSMB2* when compared to a reference standard, wherein the reference standard is from a granulocyte unsuitable for treating cancer; and/or

b. decreased expression of *ANXA1* and/or *PPP3CB* when compared to a reference standard, wherein the reference standard is from a granulocyte unsuitable for treating cancer.

23. The stem cell according to claim 21 or 22, wherein the stem cell is an induced pluripotent stem cell, or wherein the stem cell is a haematopoietic stem cell, a common myeloid progenitor cell, a myeloblast, a N. promyelocyte, a N. myelocyte, a N. metamyelocyte, a N. band, or combinations thereof; preferably wherein the stem cell is a haematopoietic stem cell.

24. A method for producing a granulocyte for treating cancer, the method comprising:

a. providing a cell; and

b. converting the cell into a granulocyte wherein:

i. the measured expression level of one or more of *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1,* and *PSMB2* is increased in the granulocyte when compared to a reference standard when the reference standard is from a granulocyte unsuitable for treating cancer; or

ii. the measured expression level of *ANXA1* and/or *PPP3CB* is decreased in the granucloyte when compared to the reference standard, when the reference standard is from a granulocyte unsuitable for treating cancer; or

iii. the measured expression level of one or more of *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1,* and *PSMB2* is increased or the same when compared to the reference standard, when the reference standard is from a granulocyte suitable for treating cancer; or

iv. the measured expression level of *ANXA1* and/or *PPP3CB* is decreased or the same when compared to the reference standard, when the reference standard is from a granulocyte suitable for treating cancer; and

optionally isolating the granulocyte.

25. The method according to claim 24 wherein the cell is a stem cell according to any one of claims 21-23.

26. The method according to claim 25 wherein converting the stem cell into a granulocyte comprises differentiating the stem cell into a granulocyte.

27. The method according to claim 24 wherein the cell is a somatic/differentiated cell, optionally from a donor who produces granulocytes suitable for treating cancer as determined according to any one of claims 2-3 or 7-13.

28. The method according to claim 27 wherein converting the somatic/differentiated cell into a granulocyte comprises transdifferentiating the somatic/differentiated cell into a granulocyte.

29. A granulocyte (e.g. obtainable by the method according to any one of claims 13, 19-20 or 24-28), wherein the granulocyte comprises:

a. increased expression of one or more of *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1,* and *PSMB2* when compared to a reference standard, wherein the reference standard is from a granulocyte unsuitable for treating cancer; and/or

b. decreased expression of *ANXA1* and/or *PPP3CB* when compared to a reference standard, wherein the reference standard is from a granulocyte unsuitable for treating cancer.

30. A composition for treating cancer (e.g. obtainable by the method according to any one of claims 13, 19-20 or 24-28), the composition comprising granulocytes: wherein at least 90% of the granulocytes comprised in the composition have:

a. increased expression of one or more of *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1,* and *PSMB2* when compared to a reference standard, wherein the reference standard is from a granulocyte unsuitable for treating cancer; and/or

b. decreased expression of *ANXA1* and/or *PPP3CB* when compared to a reference standard, wherein the reference standard is from a granulocyte unsuitable for treating cancer.

31. The composition according to claim 30, wherein at least 95%, 99% or 100% of the granulocytes comprised in the composition have:

a. increased expression of one or more of *GM2A, CTSG, CAP37, ITGB1, CYBB, SYK, DOCK8, COMP, ATG7, SLC2A1, GZMK, ATM, IKBKB, BCAP31, TAPBP, PERM, PLEC, ACSL1, RAC1,* and *PSMB2* when compared to a reference standard, wherein the reference standard is from a granulocyte unsuitable for treating cancer; and/or

b. decreased expression of *ANXA1* and/or *PPP3CB* when compared to a reference standard, wherein the reference standard is from a granulocyte unsuitable for treating cancer.

32. A pharmaceutical composition comprising:

a. the granulocyte according to claim 29, the stem cell according to any one of claims 21-23, or the composition according to claim 30 or 31; and
b. a pharmaceutically acceptable carrier, excipient, adjuvant, and/or salt.

33. The pharmaceutical composition according to claim 32, further comprising TNF-alpha.

34. The pharmaceutical composition according to any one of claims 32-33, further comprising a granulocyte-macrophage colony-stimulating factor (GM-CSF), a granulocyte colony-stimulating factor (G-CSF), a growth hormone, and serotonin, vitamin C, vitamin D, glutamine (Gln), arachidonic acid, AGE-albumin, an interleukin, TNF-alpha, Flt-3 ligand, thrombopoietin, foetal bovine serum (FBS), retinoic acid, lipopolysaccharide (LPS), IFN-gamma, IFN-beta, or combinations thereof.

35. The pharmaceutical composition according to any one of claims 32-34, further comprising a granulocyte-macrophage colony-stimulating factor (GM-CSF) and IFN-gamma.

36. A kit comprising:
a. the granulocyte according to claim 29, the stem cell according to any one of claims 21-23, the composition according to claim 30 or 31, or the pharmaceutical composition according to any one of claims 32-35; and
b. instructions for use of the same in medicine.

37. A granulocyte according to claim 29, stem cell according to any one of claims 21-23, composition according to claim 30 or 31, pharmaceutical composition according to any one of claims 32-35, or kit according to claim 36 for use in treating cancer.

38. Use of the granulocyte according to claim 29, the stem cell according to any one of claims 21-23, the composition according to claim 30 or 31, the pharmaceutical composition according to any one of claims 32-35, or the kit according to claim 36 in the manufacture of a medicament for treating cancer.

39. A method for treating cancer comprising: administering to a subject in need thereof the granulocyte according to claim 29, the stem cell according to any one of claims 21-

23, the composition according to claim 30 or 31, the pharmaceutical composition according to any one of claims 32-35, or the kit according to claim 36.

40. The granulocyte for use, stem cell for use, composition for use, pharmaceutical composition for use, or kit for use according to claim 37, the use according to claim 38 or the method according to claim 39, wherein the cancer is one or more of: pancreatic cancer, liver cancer, oesophageal cancer, stomach cancer, cervical cancer, ovarian cancer, lung cancer, bladder cancer, kidney cancer, brain cancer, prostate cancer, myeloma cancer, non-Hodgkin's lymphoma (NHL), larynx cancer, uterine cancer, or breast cancer; preferably wherein the cancer is pancreatic cancer.

41. A cell bank comprising the granulocyte according to claim 29, the stem cell according to any one of claims 21-23, the composition according to claim 30 or 31, or the pharmaceutical composition according to any one of claims 32-35.