

## CLAIMS

### WHAT IS CLAIMED IS:

1. A protein comprising:
  - (a) a first antigen-binding site that binds NKG2D;
  - (b) a second antigen-binding site that binds CD33; and
  - (c) an antibody Fc domain or a portion thereof sufficient to bind CD16.
2. The protein according to claim 1, wherein the first antigen-binding site binds to human and non-human primate NKG2D.
3. The protein according to claim 1 or 2, wherein the first antigen-binding site comprises a heavy chain variable domain and a light chain variable domain.
4. A protein according to claim 3, wherein the heavy chain variable domain and the light chain variable domain are present on the same polypeptide.
5. The protein according to any one of claims 3-4, wherein the second antigen-binding site comprises a heavy chain variable domain and a light chain variable domain.
6. The protein according to claim 5, wherein the heavy chain variable domain and the light chain variable domain of the second antigen-binding site are present on the same polypeptide.
7. The protein according to claim 5 or 6, wherein the light chain variable domain of the first antigen-binding site has an amino acid sequence identical to the amino acid sequence of the light chain variable domain of the second antigen-binding site.
8. The protein according to any one of claims 1-7, wherein the first antigen-binding site comprises
  - (a) a heavy chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:79, a heavy chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:80, a heavy chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:81, a light chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:82, a light chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:83, and a light chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:84;

- (b) a heavy chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:54, a heavy chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:55, and a heavy chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:56;
- (c) a heavy chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:57, a heavy chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:58, a heavy chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:59, a light chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:60, a light chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:61, and a light chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:62;
- (d) a heavy chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:63, a heavy chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:64, a heavy chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:65, a light chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:66, a light chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:67, and a light chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:68;
- (e) a heavy chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:71, a heavy chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:72, a heavy chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:73, a light chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:74, a light chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:75, and a light chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:76;
- (f) a heavy chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:87, a heavy chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:88, a heavy chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:89, a light chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:90, a light chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:91, and a light chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:92; or
- (g) a heavy chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:135, a heavy chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:136, a heavy chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:137, a light chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:138, a light chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:139, and a light chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:140.

9. The protein according to claim 8, wherein the first antigen-binding site comprises a heavy chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:1.
10. The protein according to claim 8, wherein the first antigen-binding site comprises a heavy chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:41 and a light chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:42.
11. The protein according to claim 8, wherein the first antigen-binding site comprises a heavy chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:43 and a light chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:44.
12. The protein according to claim 8, wherein the first antigen-binding site comprises a heavy chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:45 and a light chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:46.
13. The protein according to claim 8, wherein the first antigen-binding site comprises a heavy chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:47 and a light chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:48.
14. The protein according to claim 8, wherein the first antigen-binding site comprises a heavy chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:69 and a light chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:70.
15. The protein according to claim 8, wherein the first antigen-binding site comprises a heavy chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:77 and a light chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:78.
16. The protein according to claim 8, wherein the first antigen-binding site comprises a heavy chain variable domain amino acid sequence at least 95% identical to the amino acid

sequence of SEQ ID NO:85 and a light chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:86.

17. The protein according to claim 8, wherein the first antigen-binding site comprises a heavy chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:133 and a light chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:134.

18. The protein of claim 1 or 2, wherein the first antigen-binding site is a single-domain antibody.

19. The protein of claim 18, wherein the single-domain antibody is a V<sub>H</sub>H fragment or a V<sub>NAR</sub> fragment.

20. The protein according to any one of claims 1-2 and 18-19, wherein the second antigen-binding site comprises a heavy chain variable domain and a light chain variable domain.

21. The protein according to claim 20, wherein the heavy chain variable domain and the light chain variable domain of the second antigen-binding site are present on the same polypeptide.

22. The protein according to any one of claims 1-21, wherein the second antigen-binding site comprises

(a) a heavy chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:95, a heavy chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:96, a heavy chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:97, a light chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:98, a light chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:99, and a light chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:100;

(b) a heavy chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:103, a heavy chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:104, a heavy chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:105, a light chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:106, a light chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:107, and a light chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:108;

(c) a heavy chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:111, a heavy chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:112, a heavy chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:113, a light chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:114, a light chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:115, and a light chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:116;

(d) a heavy chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:119, a heavy chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:120, a heavy chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:121, a light chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:122, a light chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:123, and a light chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:124; or

(e) a heavy chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:127, a heavy chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:128, a heavy chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:129, a light chain CDR1 sequence identical to the amino acid sequence of SEQ ID NO:130, a light chain CDR2 sequence identical to the amino acid sequence of SEQ ID NO:131, and a light chain CDR3 sequence identical to the amino acid sequence of SEQ ID NO:132.

23. The protein according claim 22, wherein the second antigen-binding site comprises a heavy chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:93 and a light chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:94.

24. The protein according claim 22, wherein the second antigen-binding site comprises a heavy chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:101 and a light chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:102.

25. The protein according claim 22, wherein the second antigen-binding site comprises a heavy chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:109 and a light chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:110.

26. The protein according claim 22, wherein the second antigen-binding site comprises a heavy chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:117 and a light chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:118.
27. The protein according claim 22, wherein the second antigen-binding site comprises a heavy chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:125 and a light chain variable domain amino acid sequence at least 95% identical to the amino acid sequence of SEQ ID NO:126.
28. The protein according to any one of claims 1-4 and 8-17, wherein the second antigen-binding site is a single-domain antibody.
29. The protein of claim 28, wherein the second antigen-binding site is a V<sub>H</sub>H fragment or a V<sub>NAR</sub> fragment.
30. The protein according to any one of the preceding claims, wherein the protein comprises a portion of an antibody Fc domain sufficient to bind CD16, wherein the antibody Fc domain comprises hinge and CH2 domains.
31. The protein according to claim 30, wherein the antibody Fc domain comprises hinge and CH2 domains of a human IgG1 antibody.
32. The protein according to claim 30 or 31, wherein the Fc domain comprises an amino acid sequence at least 90% identical to amino acids 234-332 of a human IgG1 antibody.
33. The protein according to any one of claims 30-32, wherein the Fc domain comprises an amino acid sequence at least 90% identical to the Fc domain of human IgG1 and differs at one or more positions selected from the group consisting of Q347, Y349, T350, L351, S354, E356, E357, K360, Q362, S364, T366, L368, K370, N390, K392, T394, D399, S400, D401, F405, Y407, K409, T411, and K439.
34. A formulation comprising the protein according to any one of the preceding claims and a pharmaceutically acceptable carrier.
35. A host or *in vitro* cell comprising one or more nucleic acids encoding the protein according to any one of claims 1-33.

36. Use of the protein according to any one of claims 1-33 in the manufacture of a medicament for directly and/or indirectly enhancing CD33-expressing tumor cell death by exposing the tumor cell and a natural killer cell to an effective amount of the protein.
37. Use of the protein according to any one of claims 1-33 or the formulation according to claim 34 in the manufacture of a medicament for treating CD33-expressing cancer.
38. The use of claim 37, wherein the CD33-expressing cancer is AML, myelodysplastic syndromes, chronic myelomonocytic leukemia, myeloid blast crisis of chronic myeloid leukemia, or ALLs.