

## NUCLEIC ACID-POLYPEPTIDE COMPOSITIONS AND USES THEREOF

### CROSS-REFERENCE

[0001] This application claims the benefit of U.S. Provisional Application No. 62/316,919, filed April 1, 2016, which application is incorporated herein by reference.

### SEQUENCE LISTING

[0002] The instant application contains a Sequence Listing which has been submitted electronically in ASCII format and is hereby incorporated by reference in its entirety. Said ASCII copy, created on March 28, 2017, is named 45532-707\_601\_SL.txt and is 615,666 bytes in size.

### BACKGROUND OF THE DISCLOSURE

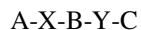
[0003] Gene suppression by RNA-induced gene silencing provides several levels of control: transcription inactivation, small interfering RNA (siRNA)-induced mRNA degradation, and siRNA-induced transcriptional attenuation. In some instances, RNA interference (RNAi) provides long lasting effect over multiple cell divisions. As such, RNAi represents a viable method useful for drug target validation, gene function analysis, pathway analysis, and disease therapeutics.

### SUMMARY OF THE DISCLOSURE

[0003a] In one aspect, disclosed herein is a composition comprising: an anti-~~CD71~~~~transferrin receptor-1~~ antibody or ~~its~~ antigen binding fragments thereof; a double-stranded siRNA molecule consisting of a passenger strand and a guide strand; and a polymer conjugated to the 3' terminus of the passenger strand; wherein said double-stranded siRNA molecule is conjugated to a cysteine of said anti-CD71 antibody or antigen binding fragments thereof by a linker; wherein said double-stranded siRNA molecule comprises at least one 2' modified nucleotide, at least one modified internucleotide linkage, or at least one inverted abasic moiety; and wherein the linker is conjugated to the 5' terminus of the passenger strand of said double stranded siRNA molecule.

[0004] Disclosed herein, in certain examples, are compositions and pharmaceutical formulations that comprise a binding moiety conjugated to a polynucleic acid molecule and a polymer. In some examples, also described herein include methods for treating a disease or condition (e.g., cancer) that utilize a composition or a pharmaceutical formulation comprising a binding moiety conjugated to a polynucleic acid molecule and a polymer.

[0005] Disclosed herein, in certain examples, is a molecule of Formula (I):



Formula I

wherein,

A is a binding moiety;

B is a polynucleotide;

C is a polymer;

## CLAIMS

### WHAT IS CLAIMED IS:

1. A composition comprising:

an anti-~~transferrin receptor-1~~CD71 antibody or ~~its~~antigen binding fragments thereof;

a double-stranded siRNA molecule consisting of a passenger strand and a guide strand; and

a polymer conjugated to the 3' terminus of the passenger strand;

wherein said double-stranded siRNA molecule is conjugated to a cysteine of said anti-CD71 antibody or antigen binding fragments thereof by a linker;

wherein said double-stranded siRNA molecule comprises at least one 2' modified nucleotide, at least one modified internucleotide linkage, or at least one inverted abasic moiety; and

wherein the linker is conjugated to the 5' terminus of the passenger strand of said double stranded siRNA molecule.

2. The composition of claim 1, wherein the linker is a cleavable linker.

3. The composition of claim 1, wherein the linker is a non-cleavable linker.

4. The composition of claim 1, wherein the polymer is polyethylene glycol (PEG).

5. The composition of claim 4, wherein the PEG has a molecular weight from about 1000 Da to about 5000 Da.

6. The composition of claim 4, wherein the PEG has a molecular weight of about 1000 Da, 2000 Da, or 5000 Da.

7. The composition of claim 1, wherein the anti-CD71 antibody or antigen binding fragment thereof comprises a humanized antibody or binding fragment thereof, chimeric antibody or binding fragment thereof, monoclonal antibody or binding fragment thereof, monovalent Fab', divalent Fab2 or single-chain variable fragment (scFv).

8. The composition of claim 6, wherein the at least one 2' modified nucleotide comprises 2'-O-methyl, 2'-O-methoxyethyl (2'-O-MOE), 2'-O-aminopropyl, 2'-

deoxy, 2'-fluoro, 2'-O-aminopropyl (2'-O-AP), 2'-O-dimethylaminoethyl (2'-O-DMAOE), 2'-O-dimethylaminopropyl (2'-O-DMAP), 2'-O-dimethylaminoethoxyethyl (2'-O-DMAEOE), or 2'-O-N-methylacetamido (2'-O-NMA) modified nucleotide.

9. The composition of claim 6, wherein the at least one 2' modified nucleotide comprises locked nucleic acid (LNA) or ethylene nucleic acid (ENA).

10. The composition of claim 6, wherein the at least one modified internucleotide linkage comprises a phosphorothioate linkage or a phosphorodithioate linkage.

11. The composition of claim 6, wherein the at least one inverted abasic moiety is at the 5' terminus ~~least one~~ or 3' terminus of the double-stranded siRNA molecule.

12. The composition of claim 6, wherein the guide strand hybridizes to a target region of a gene selected from the group consisting of KRAS, EGFR, AR, CTNNB1, PIK3CA, PIK3CB, MYC, and HPRT1.

13. The composition of claim 6, wherein the guide strand comprises a sequence selected from SEQ ID NOs: 17, 19, 21, 23, 25, 27, 29, 31, 33, 35, 37, 39, 41, 43, 45, 47, 49, 51, 53, 55, 57, 59, 61, 63, 65, 67, 69, 71, 73, 75, 453, 455, 457, 459, 461, 463, 465, 467, 469, 471, 473, 475, 477, 479, 481, 483, 485, 487, 489, 491, 493, 495, 497, 499, 501, 503, 505, 507, 509, 511, 513, 515, 517, 519, 521, 523, 525, 527, 529, 531, 533, 535, 537, 539, 541, 543, 545, 547, 549, 551, 553, 555, 557, 559, 561, 563, 565, 567, 569, 571, 573, 575, 577, 579, 581, 583, 585, 587, 589, 591, 593, 595, 597, 599, 601, 603, 605, 607, 609, 611, 613, 615, 617, 619, 621, 623, 625, 627, 629, 631, 633, 635, 637, 639, 641, 643, 645, 647, 649, 651, 653, 655, 657, 659, 661, 663, 665, 667, 669, 671, 673, 675, 677, 679, 681, 683, 685, 687, 689, 691, 693, 695, 697, 699, 701, 703, 705, 707, 709, 711, 713, 715, 717, 719, 721, 723, 725, 727, 729, 731, 733, 735, 737, 739, 741, 743, 745, 747, 749, 751, 753, 755, 757, 759, 761, 763, 765, 767, 769, 771, 773, 775, 777, 779, 781, 783, 785, 787, 789, 791, 793, 795, 797, 799, 801, 803, 805, 807, 809, 811, 813, 815, 817, 819, 821, 823, 825, 827, 829, 831, 833, 835, 837, 839, 841, 843, 845, 847, 849, 851, 853, 855, 857, 859, 861, 863, 865, 867, 869, 871, 873, 875, 877, 879, 881, 883, 885, 887, 889, 891, 893, 895, 897, 899, 901, 903, 905, 907, 909, 911, 913, 915, 917, 919, 921, 923, 925, 927, 929, 931, 933, 935, 937, 939, 941, 943, 945, 947, 949, 951, 953, 955, 957, 959, 961, 963, 965, 967, 969, 971, 973, 975, 977, 979, 981, 983, 985, 987, 989, 991, 993, 995, 997, 999, 1001, 1003, 1005, 1007, 1009, 1011, 1013, 1015, 1017, 1019, 1021, 1023, 1025, 1027, 1029, 1031, 1033, 1035, 1037, 1039, 1041, 1043, 1045, 1047, 1049, 1051, 1053, 1055, 1057, 1059, 1061, 1063, 1065, 1067, 1069, 1071, 1073, 1075, 1077, 1079, 1081,

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