CLAIMS

We claim:

1. A polynucleotide encoding a polypeptide comprising:

(a) at least one fragment of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 1,

(b) at least one fragment of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 6,

(c) at least one fragment of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 11, and

(d) at least one fragment of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 13;

wherein the polynucleotide is operatively linked to one or more sequences which direct expression of said polypeptide in a host cell;

wherein the polypeptide further comprises at least one fragment of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 21;

wherein the polypeptide comprises:

(a) a first and a second fragment of Latent Membrane Protein 1 (LMP1), wherein said first and second fragments of LMP1 are selected from the group consisting of SEQ ID NOs: 2-5 and wherein said first and second fragments of LMP1 are not adjacent to each other in the polypeptide,

(b) a first and a second fragment of Latent Membrane Protein 2 (LMP2), wherein

said first and second fragments of LMP1 are selected from the group consisting of SEQ ID NOs: 7-10 and wherein said first and second fragments of LMP2 are not adjacent to each other in the polypeptide,

(c) a fragment of Ebstein-Barr Nuclear Antigen 1 (EBNA1) consisting of SEQ ID NO: 12, and

(d) a first and a second fragment of Ebstein-Barr Nuclear Antigen 3A (EBNA3A), wherein said first and second fragments of EBNA3A are selected from the group consisting of SEQ ID NOs: 14- 20 and wherein said first and second fragments of EBNA3A are not adjacent to each other in the polypeptide.

~~2. The polynucleotide of claim 1, wherein the polypeptide further comprises at least one fragment of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 21.~~

~~3~~2. The polynucleotide of claim~~s~~ 1~~ or 2~~, wherein the polypeptide comprises:

(e) at least two fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 1,

(f) at least two fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 6,

(g) at least two fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 11,

(h) at least two fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 13, or

(i) at least two fragments of at least 8, at least 9, at least 10, at least 11, at

least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 21.

43. The polynucleotide of any of the preceding claims, wherein the polypeptide comprises:

(a) at least three fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 1,

(b) at least three fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 6,

(c) at least three fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least NO: 11,

(d) at least three fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 13, or

(e) at least three fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 21.

54. The polynucleotide of any of the preceding claims, wherein the polypeptide comprises:

(a) at least four fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 1,

(b) at least four fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 6,

(c) at least four fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 11,

(d) at least four fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 13, or

(e) at least four fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least NO: 21.

~~6~~5. The polynucleotide of any of the preceding claims, wherein the polypeptide comprises:

(a) at least five fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 1,

(b) at least five fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 6,

(c) at least five fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 11,

(d) at least five fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least

50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 13, or

(e) at least five fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 21.

~~7~~6. The polynucleotide of any of the preceding claims, wherein the polypeptide comprises:

(a) at least six fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 1,

(b) at least six fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 6,

(c) at least six fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 11,

(d) at least six fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 13, or

(e) at least six fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 21.

~~8~~7. The polynucleotide of any of the preceding claims, wherein the polypeptide comprises:

(a) at least seven fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 1,

(b) at least seven fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 6,

(c) at least seven fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 11,

(d) at least seven fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 13, or

(e) at least seven fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 21.

~~9~~8. The polynucleotide of any of the preceding claims, wherein the polypeptide comprises:

(a) at least eight fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 1,

(b) at least eight fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 6,

(c) at least eight fragments of at least 8, at least 9, at least 10, at least 11, at

least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 11,

(d) at least eight fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 13, or

(e) at least eight fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 21.

~~109~~9. The polynucleotide of any of the preceding claims, wherein the polypeptide comprises:

(a) at least nine fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 1,

(b) at least nine fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 6,

(c) at least nine fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 11,

(d) at least nine fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 13, or

(e) at least nine fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least

50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 21.

~~11~~10. The polynucleotide of any of the preceding claims, wherein the polypeptide comprises:

(a) at least ten fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 1,

(b) at least ten fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 6,

(c) at least ten fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 11,

(d) at least ten fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 13, or

(e) at least ten fragments of at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 30, at least 40, at least 50, at least 60, at least 70, at least 80 at least 90 or at least 100 amino acids of SEQ ID NO: 21.

~~12~~11. The polynucleotide of any one of claims ~~3~~2-~~11~~10, wherein the fragments of SEQ ID NO: 1 are not adjacent to each other.

~~13~~12. The polynucleotide of any one of claims ~~3~~2-~~12~~11, wherein the fragments of SEQ ID NO: 6 are not adjacent to each other.

~~14~~13. The polynucleotide of any one of claims ~~3~~2-~~13~~12, wherein the fragments of SEQ ID NO: 11 are not adjacent to each other.

~~15~~14. The polynucleotide of any one of claims ~~3~~2-~~14~~13, wherein the fragments of SEQ ID NO: 13 are not adjacent to each other.

~~16~~15. The polynucleotide of any one of claims ~~3~~2-~~15~~14, wherein the fragments of SEQ ID NO: 21 are not adjacent to each other.

~~17. **The polynucleotide of any one of claims 1-16**, wherein the polypeptide comprises:~~

~~(a) a first and a second fragment of LMP1, wherein said first and second fragments of LMP1 are selected from the group consisting SEQ ID NOs: 2-5 and wherein said first and second fragments of LMP1 are not adjacent to each other in the polypeptide,~~

~~(b) a first and a second fragment of LMP2, wherein said first and second fragments of LMP1 are selected from the group consisting SEQ ID NOs: 7-10 and wherein said first and second fragments of LMP2 are not adjacent to each other in the polypeptide,~~

~~(c) a fragment of EBNA1 consisting of SEQ ID NO: 12, and~~

~~(d) a first and a second fragment of EBNA3A, wherein said first and second fragments of EBNA3A are selected from the group consisting SEQ ID NOs: 14- 20 and wherein said first and second fragments of EBNA3A are not adjacent to each other in the polypeptide.~~

~~18~~16. **The polynucleotide of any one of claims 1-~~17~~15**, wherein the polypeptide comprises:

(a) a first fragment of LMP1 consisting of SEQ ID NO: 2,

(b) a second fragment of LMP1 consisting of SEQ ID NO: 3,

(c) a third fragment of LMP1 consisting of SEQ ID NO: 4,

(d) a fourth fragment of LMP1 consisting of SEQ ID NO: 5,

(e) a first fragment of LMP2 consisting of SEQ ID NO: 7,

(f) a second fragment of LMP2 consisting of SEQ ID NO: 8,

(g) a third fragment of LMP2 consisting of SEQ ID NO: 9,

(h) a fourth fragment of LMP2 consisting of SEQ ID NO: 10,

(i) a first fragment of EBNA1 consisting of SEQ ID NO: 12,

(j) a first fragment of EBNA3A consisting of SEQ ID NO: 14,

(k) a second fragment of EBNA3A consisting of SEQ ID NO: 15,

(l) a third fragment of EBNA3A consisting of SEQ ID NO: 16,

(m) a fourth fragment of EBNA3A consisting of SEQ ID NO: 17,

(n) a fifth fragment of EBNA3A consisting of SEQ ID NO: 18,

(o) a sixth fragment of EBNA3A consisting of SEQ ID NO: 19, and

(p) a seventh fragment of EBNA3A consisting of SEQ ID NO: 20;

wherein the first, second, third and fourth LMP1 fragments are not adjacent to each other; the first, second, third and fourth LMP2 fragments are not adjacent to each other; and the first, second, third, fourth, fifth, sixth, and seventh EBNA3A fragments are not adjacent to each other.

~~19~~17. **The polynucleotide** of claim~~s 17 or 18~~16, wherein the polypeptide further comprises:

(a) a first fragment of ZEBRA consisting of SEQ ID NO: 22, and

(b) a second fragment of ZEBRA consisting of SEQ ID NO: 23;

wherein the first and second ZEBRA fragments are not adjacent to each other.

~~20~~18. **The polynucleotide of any of the preceding claims**, wherein the polypeptide is at least 80%, at least 85%, at least 90%, at least 95% or at least 99% identical to SEQ ID NO: 24 or SEQ ID NO: 26.

~~21~~19. The polynucleotide of any of the preceding claims, wherein the fragments are immunogenic fragments.

~~22~~20. A vector comprising the polynucleotide of any of the preceding claims.

~~23~~21. The vector of claim ~~22~~20, wherein the vector is an adenovirus vector or a

vaccinia virus vector.

2422. The vector of claims ~~22~~ 20 or ~~23~~21, wherein the vector is a non-human simian adenovirus vector.

2523. The vector of claim ~~24~~22, wherein the non-human simian adenovirus vector is a chimpanzee adenovirus vector.

2624. The vector of claim ~~22~~ 20 or ~~23~~21, wherein the vector is a modified vaccinia Ankara vector.

2725. **The vector of claims ~~22~~20-~~25~~ 23** which is a ChAd155-EBV-L expression vector comprising a nucleic acid sequence at least 90%, at least 95%, at least 96%, at least 97%, at least 98% or at least 99% identical to SEQ ID NO: 50.

2826. **The vector of claims ~~22~~20-~~25~~ 23** which is a ChAd155-EBV-LLy expression vector comprising a nucleic acid sequence at least 90%, at least 95%, at least 96%, at least 97%, at least 98% or at least 99% identical to SEQ ID NO: 51.

2927. A polypeptide encoded by the polynucleotide of any of claims 1-~~21~~ 19 or the vector of any of claims ~~22~~20-~~28~~26.

3028. **The polypeptide of claim ~~29~~ 27** which is at least 80%, at least 85%, at least 90%, at least 95% or at least 99% identical to SEQ ID NO: 24 or SEQ ID NO: 26.

3129. An immunogenic composition comprising the polynucleotide of any one of claims 1-~~21~~19, the vector of any one of claims ~~22~~20-~~28~~26, or the polypeptide of any one of claims ~~29~~27 - ~~30~~28; and a pharmaceutically acceptable excipient.

3230. The immunogenic composition of claim ~~31~~29, further comprising an adjuvant.

3331. Use of the polynucleotide of any one of claims 1-~~21~~19, the vector of any one

of claims ~~22~~20-~~28~~26, the polypeptide of any one of claims ~~29~~27-~~30~~28, or the immunogenic composition of any one of claims ~~31~~29-~~32~~ 30 in the manufacture of a medicament for the treatment or prophylaxis of a disease caused by Epstein-Barr Virus infection.

~~34~~32. The polynucleotide of any one of claims 1-~~21~~19, the vector of any one of claims ~~22~~20-~~28~~26, the polypeptide of any one of claims ~~29~~27-~~30~~28, or the immunogenic composition of any one of claims ~~31~~29-~~32~~ 30 for use in the treatment or prophylaxis of a disease caused by Epstein-Barr Virus infection.

~~35~~33. A method of producing a recombinant viral particle capable of expressing an EBV antigen, comprising expressing the vector of any one of claims ~~22~~20-~~28~~ 26 in a host cell.