SEQ ID NO: 14, SEQ ID NO: 15, SEQ ID NO: 16, SEQ ID NO: 17, SEQ ID NO: 18 and SEQ ID NO: 19.

5. The method of ~~any one of~~ claim~~s~~ ~~1~~ 2, wherein the functional variant of ZO-2/TJP2 has at least 80% sequence identity with said ZO-2/TJP2.

6. The method of ~~any one of~~ claim~~s~~ ~~3~~ 4, wherein the functional variant of the Snail zinc finger transcription factor family member has at least 80% sequence identity with said Snail zinc finger transcription factor family member.

7. The method of any one of the preceding claims, wherein the first polypeptide, the second polypeptide and the candidate agent are contacted separately or simultaneously.

8. The method of any one of the preceding claims, wherein a decrease in the binding of the first polypeptide and the second polypeptide indicates that the candidate agent is useful for treating cancer.

9. The method of any one of the preceding claims, wherein a decrease in the binding of the first polypeptide and the second polypeptide indicates that the candidate agent is useful for inhibiting tumor metastasis.

10. The method of any one of the preceding claims, wherein an increase in the binding of the first polypeptide and the second polypeptide indicates