AIDA is a differentially expressed gene in human brain metastatic breast cancer.

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Metastasis to the brain is a clinical problem in patients with breast cancer¹⁻³. We mined published microarray data^{4,5} to compare primary and metastatic tumor transcriptomes for the discovery of genes associated with brain metastasis in humans with metastatic breast cancer. We found that axin interactor, dorsalization associated, encoded by AIDA, was among the genes whose expression was most quantitatively different in the brain metastases of patients with metastatic breast cancer. AIDA mRNA was present at decreased quantities in brain metastatic tissues as compared to primary tumors of the breast. Importantly, expression of AIDA in primary tumors was significantly correlated with patient post-progression survival in patients with breast cancer. Modulation of AIDA expression may be relevant to the biology by which tumor cells metastasize from the breast to the brain.

Keywords: breast cancer, metastasis, brain metastases, central nervous system metastases, axin interactor, dorsalization associated, AIDA, systems biology of breast cancer, targeted therapeutics in breast cancer.

One report described a 34% incidence of central nervous system metastases in patients treated with trastuzumab for breast cancer². More recently, the NEfERT-T clinical trial⁶ which compared administration of either neratinib or trastuzumab in conjunction with paclitaxel demonstrated that in a randomized, controlled setting, in breast cancer patients treated with neratinib, not only was the incidence of central nervous system recurrence significantly lower, the time to central nervous system metastasis was significantly delayed as compared to patients administered trastuzumab⁶. The alarmingly high rate of central nervous system metastasis described, as well as data, both anecdotal² and from a randomized, controlled setting⁶ illustrating that treatment with trastuzumab may be associated with these events demands an enhanced understanding of the transcriptional makeup of brain metastatic tissues to support identification of therapeutic targets, whether they are treatment related or not. We performed a global comparative analysis of primary and metastatic tumors in patients with brain metastatic breast cancer^{4,5}. We discovered differential and decreased expression of the gene encoding axin interactor, dorsalization associated, AIDA, in brain metastatic tissues of patients with metastatic breast cancer.

Methods

We used datasets GSE10893⁴ and GS52604⁵ for this global differential gene expression analysis of brain metastatic breast cancer in conjunction with GEO2R. GSE10893 was generated using Agilent-011521 Human 1A Microarray G4110A technology with $n=11$ primary breast tumors and $n=4$ brain metastases from patients with breast cancer; analysis was performed using platform GPL885. GSE52604 was generated using Agilent-014850 Whole Human Genome Microarray 4x44K G4112F with $n=10$ normal breast tissues and $n=35$ brain metastases from patients with breast cancer; analysis was performed using platform GPL6480. The Benjamini and Hochberg method of p -value adjustment was used for ranking of differential expression but raw p -values were used to assess statistical significance of global differential expression. Log-transformation of data was auto-detected, and the NCBI generated category of platform annotation was used. A statistical test was performed to evaluate whether AIDA gene expression was significantly different between primary tumors of the breast and brain metastases in humans with breast cancer using a two-tailed t-test. For Kaplan-Meier survival analysis, we used the Kaplan-Meier plotter online tool⁷ for correlation of AIDA mRNA expression levels in primary tumors of the breast with post-progression survival in $n=414$ breast cancer patients.

Results

We performed global comparative transcriptome analysis of metastatic and primary tumor tissues of patients with metastatic breast, as well as normal tissues of the breast using published microarray data^{4,5} to describe the transcriptional landscape of brain metastasis in human breast cancer in an unbiased fashion and for the discovery of novel therapeutic targets.

AIDA is differentially expressed in the brain metastases of patients with brain metastatic breast cancer.

Through blind, systems-level analysis of published microarray data⁴, we identified axin interactor, dorsalization associated, encoded by AIDA, as a differentially expressed gene in the brain metastatic tissues of humans with breast cancer (Table 1). When sorting each of the genes expressed in brain metastases based on significance of difference as compared to primary tumors of the breast in patients with breast cancer, AIDA ranked 81 out of 17418 total transcripts (Chart 1), equating to 99.5% differential expression. Differential expression of AIDA in the brain metastases of patients with metastatic breast cancer was statistically significant (Chart 1; $p=1.65e-04$).

Differential gene expression analyses that utilize primary and metastatic tumor tissues to discover genes associated with metastasis can be challenging due to the relatively heterogeneous cellular composition and constitution of the tumor as compared to normal, untransformed (benign) tissues. Thus, to attempt to validate differential transcriptome-wide differential expression of AIDA in brain metastases in human metastatic breast cancer, we queried a second microarray dataset⁵, here comparing normal breast tissues rather than primary tumors of the breast to brain metastases. Again, we identified AIDA as a differentially expressed gene in the brain metastatic tissues of patients with breast cancer (Chart 2). When sorting each of the genes expressed in brain metastases based on significance of difference as compared to normal breast tissues, AIDA ranked 2848 out of 41093 total transcripts (Chart 2), equating to 93.1% differential expression. Differential expression of AIDA in the brain metastases of patients with metastatic breast cancer was statistically significant (Chart 2; $p=1.88e-05$). Thus, differential expression of AIDA, transcriptome-wide, in the brain metastases of women with metastatic breast cancer was conserved across two independent microarray datasets, both when compared to benign and transformed breast tissues. This suggests that AIDA differential expression in human brain metastatic breast cancer is a *bona fide* biological process.

AIDA is expressed at lower levels in the brain metastases of patients with metastatic breast cancer.

We obtained exact mRNA expression levels for AIDA, in primary tumors of the breast and in brain metastasis of patients with brain metastatic breast cancer to determine direction and statistical significance of change in AIDA expression in brain metastatic tissues. We observed significantly decreased expression of AIDA in the brain metastases of patients with breast cancer as compared to primary tumors of the breast: AIDA was expressed at 0.24 ± 0.36 arbitrary units (AU) in primary tumors of the breast, while it was expressed at -1.03 ± 0.67 AU in brain metastatic tissues (Figure 1). The difference in AIDA mRNA levels between primary tumors of the breast and brain metastatic tissues was statistically significant (Figure 1; $p=0.000322$).

AIDA expression is correlated with survival outcomes in human breast cancer.

We performed Kaplan-Meier survival analysis⁷ in 414 breast cancer patients in total, to evaluate whether AIDA tumor expression was correlated with survival outcomes in breast cancer. We observed a correlation between primary tumor expression of AIDA and post-progression survival (PPS) in patients with breast cancer, in the upper survival quartile (Figure 2). Patients whose primary tumors expressed low levels of AIDA possessed median PPS of 34.92 months, while patients whose tumors expressed high levels of AIDA possessed median PPS of 26.04 months. This difference in PPS based on AIDA tumor expression in patients with breast cancer trended towards statistical significance (Figure 2, Chart 3; logrank p -value 0.075; hazard ratio: (0.98-1.59)).

Thus, by mining published microarray data^{4,5} in an unbiased and systematic fashion, we identified axin interactor, dorsalization associated, encoded by AIDA, as among the genes whose expression was most different, transcriptome-wide, in the brain metastases of patients with breast cancer, when compared to primary tumors of the breast and to normal breast tissues, respectively; we observed significantly decreased expression of AIDA in brain metastases as compared to primary tumors of the breast. Further, we found a correlation between AIDA expression and patient survival outcomes, as post-progression survival was greater in patients whose primary tumors expressed higher levels of AIDA as compared to patients whose primary tumors expressed lower levels of AIDA.

Discussion

We provided evidence here that AIDA is among the genes whose expression is most different in the brain metastases of patients with metastatic breast cancer, that AIDA mRNA is present at significantly decreased quantity in brain metastatic tissues as compared to primary tumors of the breast, and that primary tumor AIDA expression is correlated with patient survival outcomes in human breast cancer. Evaluation of the effects of genetic depletion of AIDA in mouse models of metastatic breast cancer on metastasis to the central nervous system is merited. Modulation of AIDA expression may be relevant to the processes by which breast cancer cells exit the breast, enter the vasculature and/or lymphatics, reside in the lymph nodes, evade immune clearance, breach the blood-brain barrier and colonize the brain.

References

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Rank: 81
Probe ID: 1985
p-value: 1.65e-04
t: 4.90
B: 1.06457
Gene: AIDA
Gene name: axin interactor, dorsalization associated

Chart 1: AIDA is differentially expressed in brain metastatic breast cancer when comparing brain metastases to primary tumors of the breast.

The rank of global differential expression, the probe/transcript ID, the *p*-value with respect to differential expression transcriptome-wide, *t*, a moderated *t*-statistic, *B*, the log-odds of differential expression between the groups compared, the gene and gene name are listed in this chart.

Rank: 2848
probe ID: A_23_P97457
p-value: 1.88e-05
t: -4.7617168
B: 2.557932
Gene: AIDA
Gene name: axin interactor, dorsalization associated

Chart 2: AIDA is differentially expressed in brain metastatic breast cancer when comparing brain metastases to normal breast tissues.

The rank of global differential expression, probe/transcript ID, the *p*-value with respect to differential expression transcriptome-wide, *t*, a moderated *t*-statistic, *B*, the log-odds of differential expression between the groups compared, the gene and gene name are listed in this chart.

Figure 1: AIDA is expressed at significantly lower levels in the brain metastases of patients with metastatic breast cancer when compared to primary tumors of the breast.

The mRNA expression level of AIDA in primary tumors of the breast (left) and in brain metastases of women with metastatic breast cancer (right) is graphically depicted; the result of a statistical test evaluating significance of difference in AIDA expression between primary tumors and brain metastases is $p=0.000322$.

Figure 2: AIDA primary tumor expression is correlated with post-progression survival in patients with breast cancer.

Depicted in this Kaplan-Meier plot is the probability of post-progression survival for $n=414$ total breast cancer patients stratified into two groups, based on low or high expression of AIDA in patient primary tumor. The log rank p -value denoting statistical significance of difference in post-progression survival when comparing the two groups, as well as hazard ratio for this comparison is listed above. Listed below is the number of patients at risk (number of patients alive) per interval, after stratification based on AIDA expression; in the first interval, number at risk is number of patients alive; in each subsequent interval, number at risk is the number at risk less those who have expired or are censored.

1 Low AIDA expression: 34.92 months

2 High AIDA expression: 26.04 months

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4 **Chart 3: Median post-progression survival is inferior in patients with high primary tumor expression of AIDA.**

5 The median PPS (post-progression survival) of patients with low primary tumor expression of AIDA and
6 high primary tumor expression of AIDA is listed in this chart, for $n=414$ breast cancer patients.