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## Suppression of the ATP-binding cassette transporter ABCC4 impairs neuroblastoma tumor growth and sensitizes to irinotecan *in vivo*

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### Abstract

The ATP-binding cassette transporter ABCC4 (Multidrug resistance protein 4, MRP4) mRNA level is a strong predictor of poor clinical outcome in neuroblastoma which may relate to its export of endogenous signaling molecules and chemotherapeutic agents. We sought to determine whether ABCC4 contributes to development, growth, and drug response in neuroblastoma *in vivo*. In neuroblastoma patients, high ABCC4 protein levels were associated with reduced overall survival.

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Inducible knockdown of ABCC4 strongly inhibited the growth of human neuroblastoma cells *in vitro* and impaired the growth of neuroblastoma xenografts. Loss of *Abcc4* in the Th-*MYCN* transgenic neuroblastoma mouse model did not impact tumor formation, however *Abcc4*-null neuroblastomas were strongly sensitized to the ABCC4 substrate drug irinotecan. Our findings demonstrate a role for ABCC4 in neuroblastoma cell proliferation and chemoresistance and provide rationale for a strategy where inhibition of ABCC4 should both attenuate the growth of neuroblastoma and sensitize tumors to ABCC4 chemotherapeutic substrates.

## Keywords

neuroblastoma; ABCC4; drug resistance; chemotherapy; xenograft

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## Introduction

Neuroblastoma is a extracranial solid tumor of infancy and early childhood arising from the sympathetic-adrenal lineage of the neural crest (1). It is a heterogeneous disease with few recurrent somatic mutations (2), and treatment for high-risk disease still relies heavily on conventional cytotoxic agents. *ABCC4*, encoding for the ATP-binding cassette transporter protein ABCC4 (Multidrug resistance protein 4, MRP4), is transcriptionally regulated by *MYCN* (3), which is a driver of neuroblastoma tumorigenesis (4) and an established poor prognostic factor (5, 6) and high *ABCC4* mRNA expression strongly predicts poor clinical outcome across multiple patient cohorts (7, 8). In cultured cells, ABCC4 confers resistance to several anti-cancer drugs, including the camptothecin irinotecan, a drug used in the treatment of neuroblastoma (9), however it is unknown whether ABCC4 protein expression has prognostic value or affects chemotherapy response in tumors. ABCC4 also exports endogenous signaling molecules that may influence tumor survival and proliferation, including cyclic nucleotides and eicosanoids (10, 11) and we previously reported that transient RNAi-mediated ABCC4 knockdown reduced proliferation and colony-forming ability in two neuroblastoma cell lines (7) in the absence of chemotherapy. More rigorous assessment of ABCC4 as a therapeutic target in neuroblastoma is clearly warranted.

Here we demonstrate that suppression of ABCC4 inhibits the growth of multiple neuroblastoma cell lines *in vitro* and established human neuroblastoma xenografts in immune-deficient mice, suggesting that blocking ABCC4 function might be beneficial in established tumors, even without chemotherapy. In a neuroblastoma-prone transgenic mouse model, constitutive absence *Abcc4* did not affect neuroblastoma formation suggesting that ABCC4 function does not contribute to the genesis of this tumor. Nonetheless, the murine neuroblastomas derived from the *Abcc4*-deficient animals were sensitized to irinotecan in an allograft model. Our findings demonstrate ABCC4 inhibition as an approach to chemosensitization of neuroblastomas.

## Materials and methods

### Tissue microarray

Tissue microarray (TMA) sections with clinical annotation, from the Children's Hospital at Westmead Tumor Bank, were stained with hematoxylin and eosin (H&E) or for ABCC4 (rat monoclonal anti-MRP4 M4I-10, Abcam ab15602, 1:50 dilution, 3 µg/mL). Cores from 98 patients diagnosed between 1979 and 2013 were scored for ABCC4 staining by a pediatric pathologist blinded to clinical parameters (12). Staining was scored for intensity (0, absent; 1, weak; 1.5 weak-moderate; 2, moderate, 2.5 moderate-strong; and 3, strong) and percentage of positive staining (0, 0%; 1, 1–10%; 2, 11–50% and 3, 51–100%) and overall score (0–9) determined by multiplying staining intensity and percentage scores, with duplicate cores averaged. Photos were from an Olympus BX53 light microscope and DP-73 camera with cellSens software.

### Cell culture

Cells lines verified by short tandem repeat profiling (CellBank Australia, Westmead Australia) were cultured at 37°C, 5% CO<sub>2</sub> in Dulbecco's Modified Eagles Medium with 10% fetal bovine serum (FBS; ThermoTrace, Nobel Park, Australia) for BE(2)-C, or Roswell Park Memorial Institute medium with 10% FBS (CHP-134 and NB69). Transfections used Lipofectamine RNAiMAX (Life Technologies, Mulgrave Australia) and 20 nM siRNA (Supplementary Table 1). Stable cell lines expressing doxycycline-inducible ABCC4 shRNA were generated by lentiviral transduction with the pFH1UTG vector (13) (Supplementary Table 1) packaged using the psPAX2 and pMD2.G plasmids (gift from Didier Trono, Lausanne, Switzerland). Doxycycline treatment was 1 µg/mL (72 h pre-treatment, media change to replenish doxycycline every 48 h).

### Western blot

Western blotting antibodies were: ABCC4 (rat monoclonal M4I-10; Enzo Life Sciences, Waterloo, NSW; 1:1000),  $\alpha$ -tubulin (mouse monoclonal DM1A; Abcam; 1:3000), total actin (rabbit polyclonal A2066; Sigma-Aldrich; 1:2000), Glyceraldehyde 3-phosphate dehydrogenase (GAPDH; mouse monoclonal G-9, sc365062; Santa Cruz Biotechnology, Dallas, TX; 1:5000), Na<sup>+</sup>/K<sup>+</sup>-ATPase  $\alpha$  (rabbit polyclonal H-300, sc-28800; Santa Cruz Biotechnology; 1:1000), peroxidase-conjugated goat anti-rat IgG (VWR International, Murarrie, QLD; 1:10000) and sheep anti-mouse horseradish peroxidase (GE Healthcare, Rydalmere, NSW; 1:50000).

### Proliferation and colony assays

Cell viability assays were in 6-well plates. Transfected cells were plated 24h after siRNA treatment, and stably transduced cells were plated in 1 µg/mL doxycycline. Viable cell number was determined using trypan blue exclusion. For colony assays, untreated or doxycycline-treated cells were plated at 200–500 cells/well, stained with 0.5% crystal violet in 50% methanol after 9–14 days, scanned, and counted using Quantity One Software (Bio-Rad) or ImageJ.

## Animal studies

All animal studies were approved by the University of New South Wales Animal Care and Ethics Committee and conducted according to the Animal Research Act, 1985 (New South Wales, Australia) and the Australian Code of Practice for Care and Use of Animals for Scientific Purposes (2013).

For xenografts,  $1 \times 10^6$  BE(2)-C cells stably transduced with inducible shRNAs were subcutaneously engrafted into one flank of Balb/c nude mice. After 7 days, mice received either control diet or diet supplemented with 600 mg/kg doxycycline (Specialty Feeds, Glen Forrest, WA, Australia). Tumors were measured every second day using a vernier caliper. Th-*MYCN* transgenic mice (4),(14) (C57BL/6, Balb/c, 129/Sv-ter background) were crossed with *Abcc4*<sup>-/-</sup> mice (C57BL/6, 129/SvJ background) (15) and Th-*MYCN*<sup>Tg/Tg</sup>/*Abcc4*<sup>+/-</sup> offspring interbred for 10 generations. Th-*MYCN*<sup>Tg/Tg</sup> mice of each *Abcc4* genotype were palpated twice weekly and humanely killed by cervical dislocation or carbon dioxide-mediated asphyxiation when a  $\sim 1000$  mm<sup>3</sup> tumor was detected.

For allograft experiments, tumors were harvested from Th-*MYCN*<sup>Tg/Tg</sup>/*Abcc4*<sup>+/+</sup> and Th-*MYCN*<sup>Tg/Tg</sup>/*Abcc4*<sup>-/-</sup> mice, dissociated ( $5 \times 10^6$  cells/mL), resuspended (1:1) with ice-cold Matrigel, and injected (0.2 mL) subcutaneously into one flank of Balb/c nude mice. Tumors were measured every second day using a vernier caliper. Drug treatment (25 mg/kg intraperitoneal (IP) irinotecan or 2 mg/kg IP cisplatin for 5 consecutive days) commenced when tumors reached 100–125 mm<sup>3</sup>. Time to reach 1000 mm<sup>3</sup> was calculated and the saline control was subtracted to account for variability between growth rates of individual tumors.

Representative tumor sections were examined tumor morphology and viability following H&E staining. Percentage of viable tumor was calculated as 100% – (% of hemorrhage, necrosis and apoptosis).

## Statistical analyses

Statistical analyses were conducted using Prism version 6 (GraphPad Software, La Jolla, CA). Survival analyses were computed by the Kaplan Meier method and compared between groups using the log-rank test. Patient overall survival (OS) was defined as the time to death within 5 years from initial diagnosis or until last contact if no event occurred. Other statistical methods are as described in the Results and figure legends.

## Results

### ABCC4 protein levels are associated with poor outcome in neuroblastoma

To investigate the relationship between ABCC4 protein expression and *MYCN* amplification and neuroblastoma outcome, tissue microarray (TMA) cores from 98 patients with primary untreated neuroblastoma (n=81), ganglioneuroma (n=12) or ganglioneuroblastoma (n=5) were immunohistochemically stained for ABCC4 (Figure 1A) and scored (0–9) for staining intensity and percentage of staining within each core. Staining of cytoplasm, neuropil, and Schwannian stroma were observed. Clinical characteristics for the cohort are shown in Table 1 and are broadly representative of the disease in terms of age at diagnosis, frequency of

*MYCN* amplification and INSS stage (16). Poorer overall survival was associated with the established prognostic indicators of INSS Stage 3 or 4 (Supplementary Figure 1A;  $P < 0.001$ ), age  $>18$  months at diagnosis (Supplementary Figure 1B;  $P = 0.047$ ) and *MYCN* amplification (Supplementary Figure 1C;  $P < 0.001$ ). *ABCC4* protein expression was higher in tumors with *MYCN* amplification (Figure 1B), with a median expression score of 6 for *MYCN*-amplified tumors and 3 for non-amplified tumors ( $< 0.001$ , Mann Whitney test). When dichotomized at the median *ABCC4* staining score ( $\leq 3$  vs  $>3$ ), the 5-year survival rates were  $74\% \pm 7\%$  and  $65\% \pm 7\%$  for patients with low or high *ABCC4* expression, respectively (Figure 1C;  $P = 0.193$ ). When dichotomized based on *ABCC4* staining score  $\leq 6$  vs  $>6$  (87 vs 11 tumors, approximately upper decile as previously described for *ABCC4* mRNA (7)), higher *ABCC4* was strongly associated with reduced overall survival ( $P = 0.014$ ), with a 5-year survival rate of  $73\% \pm 5\%$  versus  $46\% \pm 15\%$  in patients with low versus high *ABCC4* expression, respectively (Figure 1D).

### Down-regulation of *ABCC4* inhibits proliferation of human neuroblastoma cell lines

We previously showed that transient *ABCC4* knockdown (siRNAs *ABCC4.1* and *ABCC4.5*) decreased proliferation and induced neurite formation in the BE(2)-C cell line (7). To confirm that these phenotypes were attributable to *ABCC4* knockdown, we designed two additional *ABCC4* siRNA duplexes (*ABCC4.6* and *ABCC4.7*; Supplementary Table 1) and transiently transfected BE(2)-C cells. *ABCC4* protein knockdown was comparable for each duplex (Supplementary Figure 2A), as was suppression of cell proliferation in short-term assays ( $P < 0.05$ ; Supplementary Figure 2B), confirming that reduced proliferation is a consequence of *ABCC4* suppression. The percentage of neurite-positive cells was increased with *ABCC4* siRNA duplex *ABCC4.5* ( $P < 0.001$ , Supplementary Figure 2C), but not with duplexes *ABCC4.6* ( $P = 0.967$ ) or *ABCC4.7* ( $P = 0.557$ ), indicating that neurite formation is not consistently related to *ABCC4* suppression. Duplexes *ABCC4.6* and *ABCC4.7* were subsequently selected for *in vivo* studies.

To enable sustained *ABCC4* suppression, shRNA sequences equivalent to *ABCC4.6* and *ABCC4.7* and an additional shRNA targeting the *ABCC4* 3' UTR (*ABCC4.13*) were cloned into a doxycycline-inducible expression system (Supplementary Table 1). Efficient knockdown of *ABCC4* protein in transduced BE(2)-C cells was confirmed following induction with doxycycline (Figure 2A) and slowed cell proliferation in short term assays ( $>50\%$  reduction in cell number for each duplex compared to untreated cells;  $P \leq 0.05$  for each; Figure 2B, Supplementary Figure 3A–D) and in colony assays ( $>80\%$  reduction in colony numbers for each duplex;  $P \leq 0.001$ ; Figure 2C–D).

These findings were confirmed in the *MYCN*-amplified CHP-134 cell line (17), with cell numbers at 96h post-transfection 75% ( $P \leq 0.001$ ) and 62% of control ( $P \leq 0.001$ ), respectively for *ABCC4.6* and *ABCC4.7* siRNAs (Figure 3A–B) and in the non-amplified NB69 (17) cell line, with cell numbers of 55% of control ( $P \leq 0.001$ ) for *ABCC4.6* and abolition of proliferation for *ABCC4.7* (Figure 3C–D).

### ABCC4 knockdown slows the growth of xenografted human neuroblastoma cells

To test the effects of ABCC4 knockdown *in vivo*, BE(2)-C cells with stable, doxycycline-inducible ABCC4 knockdown (ABCC4.6 or ABCC4.7) were xenografted subcutaneously into Balb/c nude mice, with randomization to doxycycline or control diet 7 days later. ABCC4 knockdown increased the median time to 1000 mm<sup>3</sup> tumor from 31 to 40 days (Figure 4A) in the ABCC4.6 xenografts, and from 29 to 35 days (Figure 4B) in the ABCC4.7 xenografts. Growth curves for individual ABCC4.6 tumors are shown in Figure 4C. ABCC4 suppression was maintained in tumors harvested at ~1000 mm<sup>3</sup> (Figure 4D) albeit with a lower differential than achieved *in vitro*. On histological examination (Figure 4E), no difference in percentage viable tumor was observed between the control and doxycycline-treated tumors for either shRNA construct (64.4±13.7% and 63.5±11.0%, P = 0.961 for ABCC4.6; 79.9±7.0% and 81.1±5.7%, P = 0.892 for ABCC4.7).

### Constitutive absence of ABCC4 does not alter tumorigenesis in a murine model of neuroblastoma

To investigate whether loss of ABCC4 impacts on tumor initiation, we crossed neuroblastoma-prone Th-*MYCN* mice and mice with targeted disruption of *Abcc4*. Tumors arising in *Abcc4*<sup>+/-</sup> mice had reduced ABCC4 protein levels and those arising in *Abcc4*<sup>-/-</sup> mice had no detectable ABCC4 protein (Figure 5A). Tumor incidence was not altered by loss of either one (OR = 0.842, 95% CI = 0.516–1.38, P = 0.525, Fisher's exact test) or both *Abcc4* alleles (OR = 0.925, 95% CI = 0.525–1.63, P = 0.885, Fisher's exact test) (Figure 5B). Tumor latency was unchanged between *Abcc4* genotypes, with a median age to ~1000 mm<sup>3</sup> tumor of 81 days for *Abcc4*<sup>+/+</sup> mice, 82 days for *Abcc4*<sup>+/-</sup> mice (HR = 0.944, 95% CI = 0.753–1.18; P = 0.611) and 83 days for *Abcc4*<sup>-/-</sup> mice (HR = 0.983, 95% CI = 0.754–1.28; P = 0.895) (Figure 5C). All tumours examined were Schwannian-stroma poor neuroblastic tumours with similar degree of haemorrhage and necrosis, with no histological differences observed between genotypes (Figure 5D).

### Loss of ABCC4 alters tumor sensitivity to chemotherapeutic drugs

Next, we investigated whether ABCC4 contributes to tumor drug sensitivity in the Th-*MYCN* mouse model treated with the ABCC4 substrate irinotecan and with cisplatin, which is not an ABCC4 substrate. To avoid pharmacokinetic differences due to systemic ABCC4 loss, we harvested and dissociated tumors from Th-*MYCN*<sup>Tg/Tg</sup>/*Abcc4*<sup>+/+</sup> and Th-*MYCN*<sup>Tg/Tg</sup>/*Abcc4*<sup>-/-</sup> mice and established allografts in Balb/c nude mice, as described previously (18). Mice engrafted with *Abcc4*<sup>+/+</sup> tumors and treated with irinotecan had a median survival time of 137 days, with 2 of 10 mice surviving long-term. Survival was extended in mice engrafted with *Abcc4*<sup>-/-</sup> tumors, with 6 of 10 mice treated with irinotecan surviving long term (HR = 3.209, 95% CI = 1.081–11.07) (Figure 5E). In contrast, loss of *Abcc4*<sup>-/-</sup> did not significantly alter sensitivity to cisplatin (HR = 1.789, 95% CI = 0.769–4.887) (Figure 5F).

## Discussion

Previous studies suggest ABCC4 as an attractive therapeutic target, as its knockdown suppresses proliferation of a range of cancer cell lines *in vitro* and in xenograft models (19–

21). ABCC4 may be particularly relevant in neuroblastoma, given the association between high *ABCC4* expression and poor clinical outcome (7, 8). In contrast to previous studies which used constitutive ABCC4 knockdown, the present study mimics pharmacological inhibition by inducing knockdown in established tumours. The striking impact on cells in cell culture and the reduced growth of xenografted tumors support a role for this transporter in neuroblastoma biology.

The observation that loss of *Abcc4* has no impact on tumor formation in the Th-*MYCN* transgenic neuroblastoma mouse model has several possible non-mutually exclusive explanations. First, ABCC4 may not contribute to the genesis of neuroblastoma, but may play a role in sustaining established tumors, as indicated by the xenograft model. Second, tumors arising in *Abcc4*-null mice might harbor adaptive changes that allow their development in the absence of *Abcc4*. Third, functional differences might exist between the human and mouse protein, as suggested by differences in the effects of ABCC4 inhibition or down-regulation on human and mouse skin dendritic cell migration (22),(23) and by differences in the affinity of human and mouse ABCC4 for the substrate cGMP (24).

Notably, we observed that loss of *Abcc4* sensitizes allografted Th-*MYCN* mouse tumors to irinotecan, which along with its active metabolite SN-38 is an ABCC4 substrate (8, 25). As these experiments were conducted as allografts, this chemosensitization can be attributed to loss of tumor ABCC4 rather than to altered drug clearance as has been previously reported for another camptothecin, topotecan, in *Abcc4*-deficient mice (15). As irinotecan is used to treat relapsed neuroblastoma (9), pharmacological inhibition of ABCC4 may be of clinical benefit in this disease. Currently, no selective inhibitors of ABCC4 with proven *in vivo* efficacy are available and the most widely used small molecule inhibitor of ABCC4, the cysteinyl leukotriene receptor antagonist MK-571, also has inhibitory activity against a range of ABC transporters, including ABCC1, ABCC2, ABCC3 and ABCC5 (26, 27) as well as apparent phosphodiesterase inhibitory activity (28). While more selective ABCC4 inhibitors have been described (29), their suitability for *in vivo* use is unknown. This study provides support for the continued development of ABCC4 inhibitors, and illustrates a model in which these inhibitors could be assessed.

In conclusion, we have shown that downregulation of ABCC4 slows the progression of neuroblastoma xenografts and that loss of ABCC4 sensitizes neuroblastoma tumors to irinotecan. Inhibition of ABCC4 is therefore a potential strategy for chemosensitization in this disease, providing rationale for the further development of small molecule ABCC4 inhibitors.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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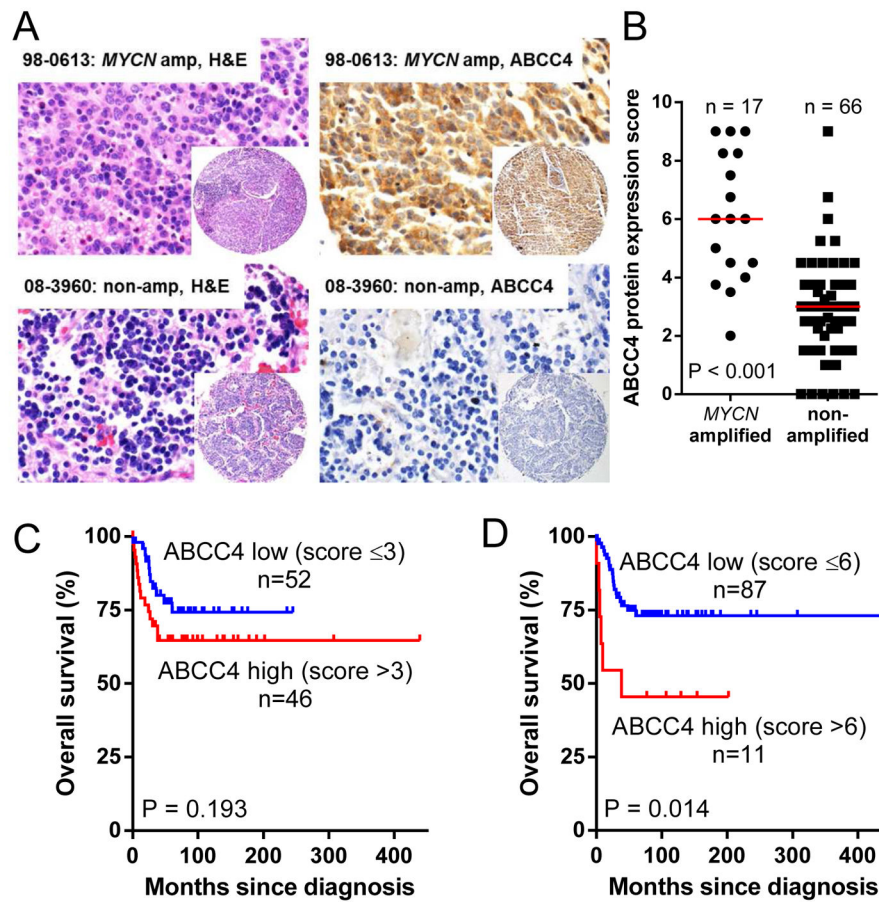
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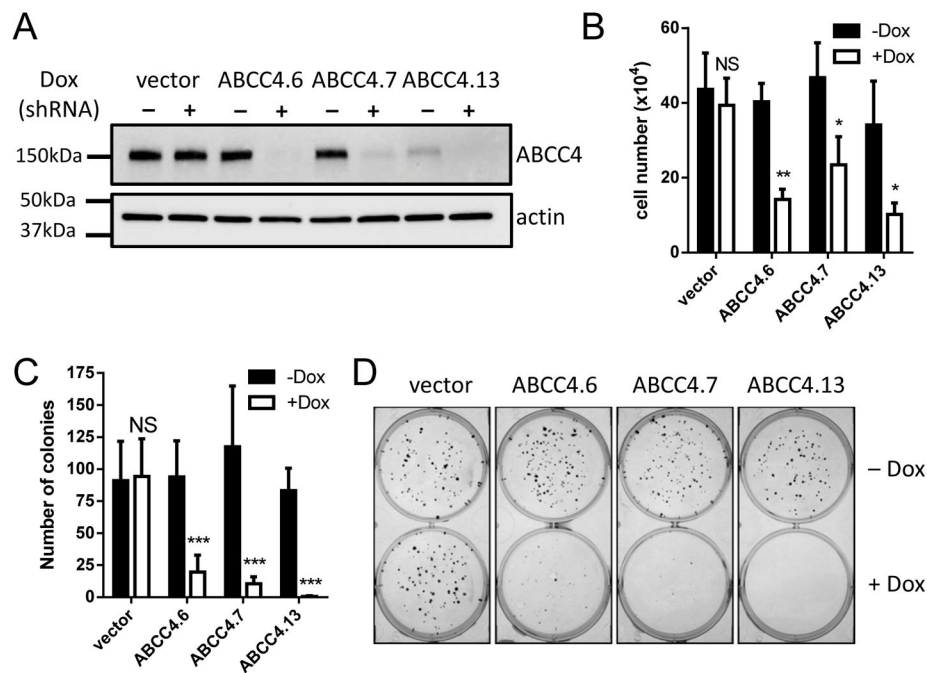
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- Suppression of ABCC4 slows progression of neuroblastoma xenografts
- Loss of ABCC4 sensitizes tumors to the camptothecin irinotecan
- Inhibition of ABCC4 may be a chemosensitization strategy to pediatric neuroblastoma



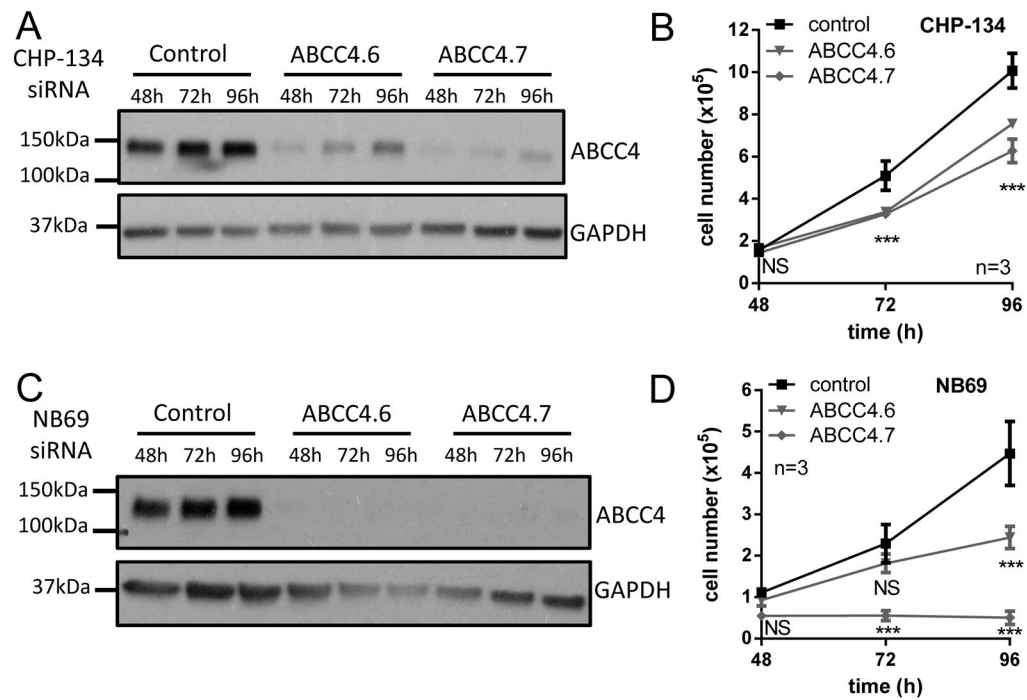
**Figure 1. ABCC4 protein levels are associated with poor outcome in neuroblastoma**

(A) Immunohistochemical staining of sample cores from a neuroblastoma TMA showing tumor morphology by H&E staining (left panels) and ABCC4 staining (right panels). A representative *MYCN*-amplified sample with high ABCC4 staining (score = 9) is shown in the upper right panel and a representative *MYCN* non-amplified, ABCC4-negative sample (score = 0) in the bottom right panel. 600x magnification for larger panels; 100x for insets. (B) Distribution of ABCC4 protein expression scores across the TMA. Bars represent median expression score, P value from Mann Whitney test. (C, D) Kaplan-Meier curves showing the probability of overall survival for patients with tumors dichotomized into high or low ABCC4 protein expression at different cut-points.



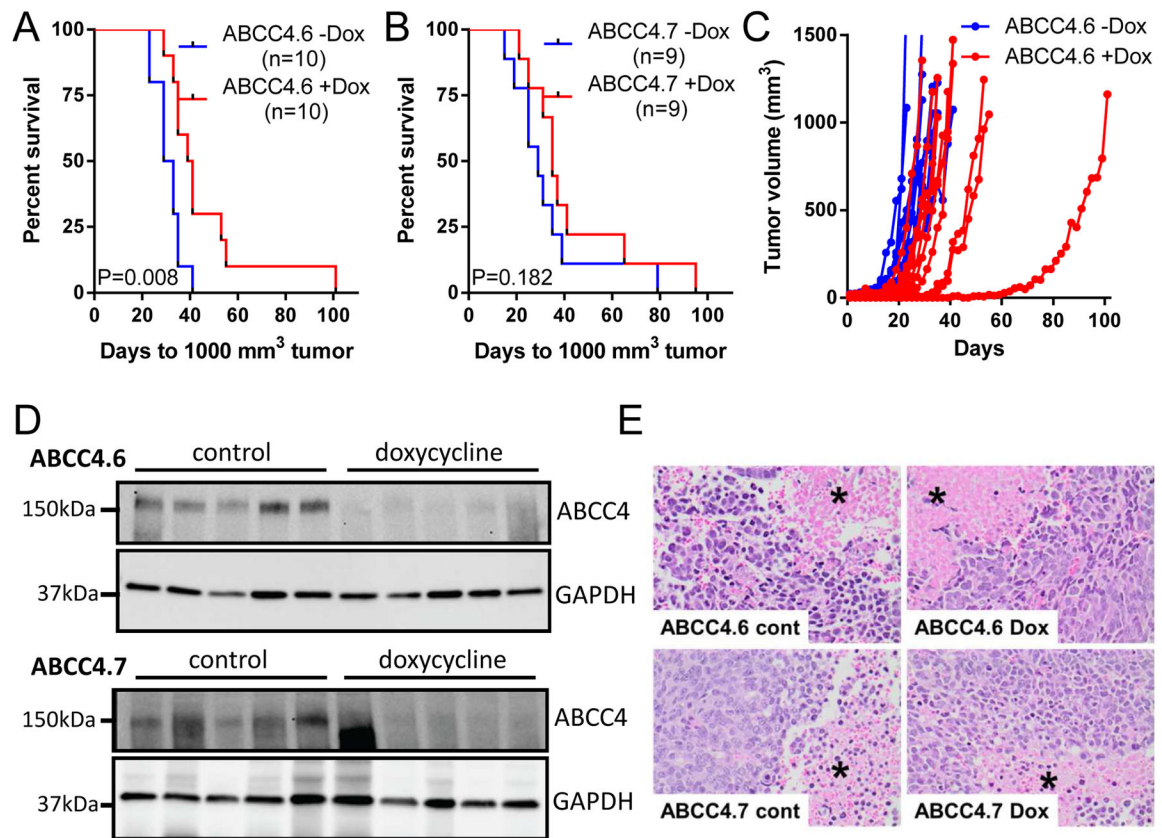
**Figure 2. Sustained suppression of ABCC4 inhibits neuroblastoma cell growth and abolishes colony formation**

(A) Efficient inducible knockdown of ABCC4 is achieved in BE(2)-C cells following doxycycline (Dox) treatment, with each of three independent shRNA duplexes (72h timepoint). Actin is shown as a loading control. (B) Inducible ABCC4 knockdown inhibits BE(2)-C cell proliferation in short-term assays at 72h post-induction for each duplex compared its untreated control (t test). (C) Inducible ABCC4 knockdown abolishes the formation of BE(2)-C cell colonies for each duplex compared to its untreated control (t test). (D) Representative images for colony assays quantitated in panel C. Data in B and C show means and SD derived from three independent experiments. \*  $P < 0.05$ ; \*\*  $P < 0.01$ ; \*\*\*  $P < 0.001$ .

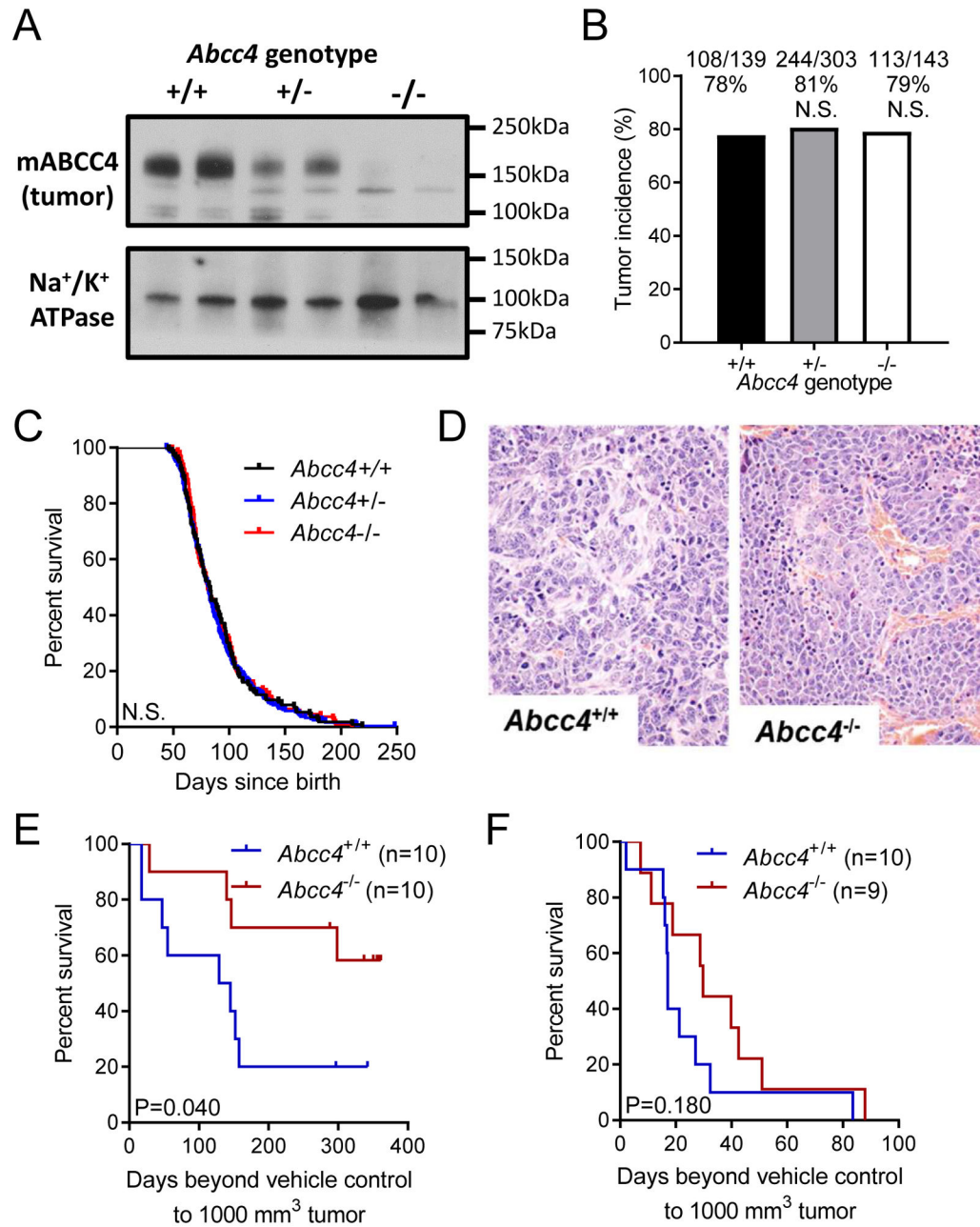


**Figure 3. Downregulation of ABCC4 inhibits proliferation of neuroblastoma cell lines**

Knockdown of ABCC4 with two independent siRNA duplexes inhibits proliferation of the *MYCN*-amplified CHP-134 cell line (A, B) and the *MYCN* single copy NB69 (C, D) cell line compared to non-silencing control siRNA at 72h. GAPDH is shown as a loading control in A and C. Data in B and D show means and SD derived from three independent experiments. \*\*\*  $P < 0.001$ , Dunnett's multiple comparisons test following 2-way ANOVA.



**Figure 4. ABCC4 knockdown slows the growth of xenografted human neuroblastoma cells**  
 (A, B) Induction of ABCC4 knockdown in established BE(2)-C tumors (doxycycline treatment commenced 7 days post-enugraftment) extends survival ( $P = 0.008$  for ABCC4.6,  $P = 0.182$  for ABCC4.7,  $n = 9-10$ /group). (C) Tumor growth in individual mice for ABCC4.6 with and without doxycycline. (D) Doxycycline-induced knockdown of ABCC4 is maintained in xenografted tumors to the experimental end-point for both ABCC4.6 and ABCC4.7. GAPDH is shown as a loading control and a different tumor sample is shown in each lane. (E) Histological appearance of representative xenografts from each cohort (H&E staining). Coagulative necrosis, where cell outlines are preserved (\*), is seen in all treatment groups. Magnification: 400x.



**Figure 5. Loss of ABCC4 does not alter tumorigenesis or tumor progression in a transgenic mouse model of neuroblastoma, but sensitizes tumors to irinotecan**  
 (A) ABCC4 protein expression in Th-*MYCN* mouse tumors derived from animals with *Abcc4*<sup>+/+</sup>, *Abcc4*<sup>+/-</sup>, and *Abcc4*<sup>-/-</sup> genotypes. Samples are membrane preparations and Na<sup>+</sup>/K<sup>+</sup> ATPase is shown as a loading control. (B) Loss of *Abcc4* does not alter tumor incidence in Th-*MYCN* mice. Number of tumor-bearing mice and total mice for each genotype are indicated above each bar (P=1.0 and P=0.725 for loss of one and both *Abcc4* alleles respectively, compared to wild-type control). (C) Loss of *Abcc4* does not alter tumor latency in Th-*MYCN* mice, as measured by median age to occurrence of a medium palpable tumor (P = 0.689 and P = 0.7071 for loss of one and both *Abcc4* alleles respectively,

compared to wild-type control). (D) Histological appearance of representative *Abcc4*<sup>+/+</sup> and *Abcc4*<sup>-/-</sup> Th-*MYCN* mouse tumors (H&E staining). Magnification: 400x. (E,F) Loss of ABCC4 significantly sensitizes allografted transgenic mouse neuroblastomas to the ABCC4 substrate irinotecan (E; P = 0.040, log-rank test), but not to cisplatin (F; P = 0.1797, log-rank test).

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**Table 1**

Clinical characteristics for the 98 patient tissue microarray cohort

Clinical characteristic	Total	%
<b>Age at diagnosis</b>		
18 months	52	53.1
> 18 months	46	46.9
<b>MYCN status</b>		
Single copy	66	67.4
Amplified	17	17.3
Unknown	15	15.3
<b>INSS stage</b>		
Stage 1,2,4S	47	48.0
Stage 3,4	50	51.0
Unknown	1	1.0
<b>Death from disease<sup>1</sup></b>	26	26.5

<sup>1</sup>Deaths attributed to neuroblastoma within 5 years of diagnosis.