Biphasic Hyalinizing Psammomatous Renal Cell Carcinoma (BHP RCC) A Distinctive Neoplasm Associated With Somatic NF2 Mutations

Pedram Argani, MD. Am J Surg Pathol. Volume 44, Number 7, July 2020

汇报人:魏洁

- Perhaps more than any other cancer, distinctive morphologic features of renal cell carcinomas (RCCs) correlate with specific genetic alterations.
 - ✓ fumarate hydratase deficiency: prominent nucleoli with a perinucleolar halo



Am J Surg Pathol. Author manuscript; available in PMC 2014 January 01.

- Perhaps more than any other cancer, distinctive morphologic features of renal cell carcinomas (RCCs) correlate with specific genetic alterations.
 - ✓ succinate dehydrogenase deficiency:solid or focally cystic growth, uniform cytology with eosinophilic flocculent cytoplasm, intracytoplasmic vacuolations and inclusions, and round to oval low-grade nuclei.



Am J Surg Pathol • Volume 38, Number 12, December 2014

- Perhaps more than any other cancer, distinctive morphologic features of renal cell carcinomas (RCCs) correlate with specific genetic alterations.
 - tuberous sclerosis complex (TSC) gene mutations: These neoplasms
 typically demonstrate solid and cystic architecture, and the neoplastic cells
 contain voluminous eosinophilic cytoplasm with granular cytoplasmic stippling .



- Perhaps more than any other cancer, distinctive morphologic features of renal cell carcinomas (RCCs) correlate with specific genetic alterations.
 - ✓ translocations involving chromosome Xp11 resulting in gene fusions involving the TFE3 transcription factor gene: the constellation of clear cells with papillary architecture, voluminous cytoplasm.



MATERIALS AND METHODS

•Cases

The 8 cases reported herein were retrieved from the files of 3 authors (P.A., V.E.R., J.N.E.) during a review of unclassified primary renal carcinomas.

Immunohistochemistry

HMB45, cathepsin K, AE1/3, Cam5.2, EMA, PAX8, S100, melan A, desmin, SF-1, GATA3, SMA, TFE3

•DNA and RNA Sequencing

•Photomicrographs

IABL	E T. Clinic	cal, Pathologic, and	d Genetic Featu	res of Cases			
Case #	Age/Sex	Tumor Size/Stage	IHC ⁺	IHC ⁻	Comment	NF2 Variant	Other Molecular Findings
1	78/male	4.2 cm; pT1bNX	HNF1-beta, PAX8, CK7, EMA (focal)	WT-1, Melan A, cathepsin k ⁻ , GATA3, SF-1, inhibin; FH, SDHB intact	TFE3/B FISH ⁻ ; died of complications 1 mo after operation	p.Tyr66*	Chr loss: 1p, 6, 22q; Chr gain: 16, 20 <i>XPO1</i> p.Asn735Lys; <i>POLQ</i> p. Phe944Val; <i>SPTBN1</i> p. Trp1893*; <i>EXT2</i> p.Tyr421*; <i>NPM1</i> p.Asp165_Glu169del; <i>NCOA1</i> p.Met476Ile; <i>AUTS2</i> p.Gly216Arg; <i>ARID2</i> p. Arg314Ser; <i>LMO7</i> p. Pro715Ser; <i>CYLD</i> p. Pro78Thr; <i>MIB1</i> p. Ser408Asn; <i>CSNK1G2</i> c.187 +1G > T
2	60/U	7 cm; pT2NX	PAX8, EMA, CK7	WT-1, TFE3, cathepsin k, TFEB, HMB45, Melan A chromogranin, synaptophysin, GATA3, SF-1, inhibin	TFE3/B FISH ⁻ ; ESRD; tumor in native kidney 21 y after transplant	p.Asn293fs	Chr loss: 1p, 6, 22q; Chr gain: 16, 20 WNT4 p.Val97Ile; RAD54L p. Arg202Cys; MRE11A p. Asp495His; ERBB2 p. Asp880Tyr; PI4KA p. Pro514Leu
3	69/male	4 cm; pT1cNXM1 (bone metastasis)	CK7, racemase, HNF1-beta	TFE3, HMB45, Melan A, cathepsin K, AR, ER, CA-IX, GATA3	Lung metastasis; died of disease 2 mo later	p.Gln111*	Chr loss: 9; Chr gain: 20 GLII p.Gly1097fs; RBI p. Glu539_Thr543del; TERT promoter variant g.1295228C > T

TABLE 1. Cl	inical, Pathologia	, and Genetic	Features of	of Cases
-------------	--------------------	---------------	-------------	----------

Case #	Ago/Soy	Tumor Size/Stage	IHC ⁺	IHC ⁻	Comment	NF2 Variant	Other Molecular Findings
4	53/female	2.5 cm; pT1NX		GATA3, WT-1, cathepsin K, inhibin, SF-1	TFE3/B FISH ⁻ ; h/o bilateral ovarian cystectomy	c.600-1G > A	Chr loss: 1p, 4, 6, 21q, 22q; Chr gain: 1, 7, 20 ARID1A p.Pro1619fs; CDK6 p. Thr198fs; RAD50 p. Gln479Arg; MLLT6 p. Ser608Tyr; SUGP2 p. Ala75Ser; THBS1 p. Arg458Gly; LRIG3 p. Asp56Glu; PLCG2 p. Glu567Lys; KMT2B p. Gly100Asp
5	73/male	3 cm; pT1aNX	CK7	CA-IX, cathepsin K, GATA3, WT-1, inhibin, SF-1 ⁻ ; FH, SDHB intact	TFE3/B FISH	p.Lys364*	Chr loss: 1, 6, 9q, 15q, 19q; Chr gain: 9p, 16 CHD6 p.Asn2152Ser; NUP98 p. Ser103*; SAMD9 p.Phe801fs; MIB1 c.2211+1G > A; CIITA p.Gly174Arg; MECOM p. Glu81Gln; FLT4 p. Trp1233Cys; ABL2 p. Asn152fs; ASPSCR1 p. Asp217fs; HIST1H2AG p. Pro81Arg; FBXW7 p. Arg505His; KDM1A p. Val400Ile; SQSTM1 p. Cys27Ser
6	71/male	1.5 cm; pT3NX (perirenal fat and vascular invasion)	PAX8, CK7	Cathepsin k, GATA3, WT-1, inhibin, SF-1 ⁻ ; FH intact	Concurrent clear cell RCC; TFE3/ B FISH ⁻	p.Leu316fs	Chr loss: 1p, 6, 9, 15q; Chr gain: 16,17q,20 SMARCA4 c.3546+3A > T; FANCD2 p.Ser352Leu; BCL2 p.Leu185Met; PAK5 p. Ala120Val; PPP2R1B p. His661Leu; NOTCH2 p. Phe2091Ser; PICALM p. Val183Gly; HIP1 c.2891- 2A > G; AKAP9 p. Glu2697Gln; STAT1 p. Ala531Thr; DLEC1 p. Gln1740Arg; PRICKLE1 p. Pro786Thr; LAMA5 p. Asn479Ser

Case #	Age/Sex	Tumor Size/Stage	ІНС⁺	IHC ⁻	Comment	NF2 Variant	Other Molecular Findings
7	52/male	1 cm; pT1NX	PAX8	Chromogranin, synaptophysin, cathepsin K, Melan A, WT-1, inhibin, SF-1, calretinin, GATA3	Renal-adrenal fusion; concurrent renal cyst	p. Arg516_Lys523- delinsGln	Chr loss: 14q, 19q, 22q; Chr gain: 2p, 5, 13q, 20 <i>PPP2R2B</i> p.Arg129fs; <i>RPL22</i> p. Lys13fs; <i>BICC1</i> p.Val254Leu; <i>WDFY3</i> c.1591+4A > C; <i>TFEB</i> p.Arg4Pro; <i>FZD3</i> p. Gly595Ala; <i>TNC</i> p. Phe1196Tyr
8	51/male	1.5 cm; pT1NX			Concurrent renal cyst	p.Lys130fs	Chr loss: 1p, 3, 6, 11, 14q, 19, 22q; Chr gain: 20 <i>ERCC5</i> p.Leu844*; <i>CLTC</i> p. Asp662fs; <i>WHSC1</i> c.1674 +6dupT; <i>SNX29</i> p.Lys513Ile; <i>CNTN1</i> c.2185-4T > A; <i>SGK1</i> p.Lys338Glu; <i>ITGA7</i> p. Glu120Val; <i>GNAI1</i> p. Arg142Ile; <i>TPR</i> p. Glu1594Asp; <i>RASGRF1</i> p. Asn1075Tyr; <i>LIFR</i> p. Met773Leu

*Stop codon.

Chr indicates chromosome; FH, fumarate hydratase; FISH, fluorescence in situ hybridization; SDHB, succinate dehydrogenase subunit B; U, unknown sex.

age: 51-78y F:M=6:1 5 underwent partial nephrectomy 2 underwent radical nephrectomy size:1-7cm; solid



FIGURE 1. BHP RCC case #3. The 4 cm neoplasm has a white, fibrous appearance, which contrasts with the yellow perinephric fat lobules to the left and the maroon-red native renal parenchyma to the upper right.



FIGURE 2. BHP RCC case #1. The neoplasm is unencapsulated and has a rounded, nodular border with the native kidney at the top of the figure (A). The biphasic pattern is evident, as smaller cells with condensed chromatin cluster around hyalinized material while larger cells with vesicular chromatin form tubules and larger acini (B).



The smaller cells also form solid spindle cell foci unassociated with basement membrane material (C). The small cells on papillae associated with the basement membrane branch within larger acini, resulting in a glomeruloid pattern (D–F).



The hyalinized material labels with type IV collagen, consistent with basement membrane material (G). The neoplasm has a low proliferative index as measured by Ki-67 IHC, particularly in the smaller cells (H).



Cytokeratin 7 preferentially labels the larger cells forming the larger acini (E). EMA preferentially labels the smaller cells in the glomeruloid bodies (F). case 2



FIGURE 4. BHP RCC case #3. The neoplasm is unencapsulated; note the native renal parenchyma at the upper left. The neoplasm has a striking biphasic appearance, with solid clusters of larger epithelioid clear cells and smaller cells forming branching papillae(A).



At higher power, one can appreciate the smaller cells with condensed chromatin clustered around basement membrane material, the larger cells with vesicular chromatin, as well as extensive psammomatous calcification (B, C).



In other areas, the neoplasm has more fibrotic stroma and the architecture is more solid with epithelioid cells surrounding smaller cells around hyalinized material and psammomatous calcifications (D).



FIGURE 5. BHP RCC case #4. At lower power, one can appreciate the unencapsulated nature of the neoplasm that borders the native kidney at the right (A). At intermediate power, one can appreciate both the tubulopapillary pattern at the right and the solid tubular pattern at the left (B).



The tubulopapillary pattern demonstrates smaller cells clustered around basement membrane material in larger acini, yielding a glomeruloid pattern (C). The more solid areas feature sclerotic stroma in which there are tubules, more cribriform structures clustered around basement membrane material, and psammomatous calcification (D).



FIGURE 6. BHP RCC case #5. Much of this neoplasm had the appearance of an unclassifiable RCC. At low power, the neoplasm demonstrates hyalinization with prominent psammomatous calcifications to the right, and a clear cell appearance to the left (A). A higher power view of the clear cell area reveals a nondescript solid clear cell proliferation that would be difficult to distinguish from high-grade conventional clear cell RCC (B).





Other areas of the neoplasm demonstrate an anastomosing tubular pattern in the myxoid stroma that is reminiscent of MTSC (note the psammomatous calcification to the right of both figures) (C, D).



In other areas, one can appreciate tubular architecture and biphasic cytology (E). Higher power view of these areas reveals larger epithelioid cells with open chromatin and eosinophilic cytoplasm and smaller cells with condensed chromatin and minimal basophilic cytoplasm (F).



FIGURE 7. BHP RCC case #6. The neoplasm demonstrated an unencapsulated border with the native kidney to the right (A). The majority of the neoplasm demonstrates the typical biphasic appearance with small cells forming glomeruloid bodies within larger acini lined by larger cells (B).



Centrally, the neoplasm demonstrates extensive sclerosis (C). Neoplastic epithelium within this sclerotic and desmoplastic stroma has a cord-like and tubular appearance that raises the differential diagnosis of collecting duct carcinoma (D).



The more typical biphasic areas (left bottom) merge with areas having a more prominent eosinophilic cytoplasm (upper right) (E). The neoplastic cells in the latter areas have abundant eosinophilic cytoplasm, vesicular chromatin, and prominent nucleoli and are associated with necrosis (F).



FIGURE 8. BHP RCC case #7. This neoplasm has an unencapsulated border with the native kidney. The dominant appearance is that of a solid tubular lesion with extensive psammomatous calcifications (A). The neoplasm intermingles among native renal elements at its border to the left (B).



At higher power, one can appreciate solid nests of neoplastic cells with vesicular chromatin, prominent nucleoli and abundant eosinophilic cytoplasm forming solid nests, frequently associated with psammomatous calcification (C, D).



FIGURE 9. BHP RCC case #8. At low power, the neoplasm has a solid and papillary architecture and has an unencapsulated border with the native kidney to the right (A). The neoplasm has a biphasic appearance, with solid, tubular, and papillary areas to the right and a nested clear cell area to the left (B).



The solid papillary areas demonstrate the typical biphasic appearance with smaller cells clustered on basement membrane material and larger cells at the periphery (C). The solid clear cell areas are indistinguishable from clear cell RCC (D).



Other areas of this neoplasm demonstrated sclerotic and desmoplastic stroma, as cords and tubules of neoplastic cells permeate among renal native tubules and glomeruli (E). In other areas, the neoplastic cells form linear cords in a desmoplastic and sclerotic stroma (F).

TABLE 2.	Morphologic	Features of	Cases

Case #	Slides	Capsule	Vaguely Nodular	Biphasic	Basement Membrane Nodules	Small Cell Spindling	Papillary	Glomeruloid Bodies	Sclerotic Stroma	Psammoma Bodies	Other
1	3	-/+	+	+	+	+	+	+	+	+	
2	3	_	+	+	+	+	+	+	+	+	Necrosis
3	1	NA	+	+	+	+	+	+	+	+	1.0010010
4	2	_*	+	+	+	+	+	+	+	+++	Necrosis
5	3	+	+	+	Minimal	+	+	-	+	+++	Clear cells, pink cells, MTSC-like
6	2	-/+*	+	Minimal	+	Minimal	+	+	+	+	Pink cells, entraps tubules
7	4	-/+	+	+	++	-	-	-	+	+++	All tubular
8	2	-	+	+	++	-	+	+	+	+	Clear cells, entraps tubules

*-prominent peritumoral chronic inflammation.
+ indicates present; ++, well-developed; +++, prominent; NA, not available.



FIGURE 10. Recurrent molecular findings in BHP RCC. A, Lollipop plot of the NF2 gene (ENST00000338641, NM_000268) showing the distribution of alterations identified. Each mutation is plotted with the corresponding case number. In-frame deletion—green; splice acceptor variant—black; frameshift or nonsense mutation—magenta. B, CNV scatter plot of case #2 showing broad copy number alterations in chromosomes 1p, 6, 16, 20, and 22q.

- ・NF2基因:
 - ✓ 肿瘤抑制基因
 - ✓ 定位与22q12.2,由17个外显子组成
 - ✓ 编码蛋白Merlin, 维持细胞骨架的稳定性, 在细胞接触时 抑制细胞增殖。
 - ✓ NF2基因还可抑制Ras所介导的肿瘤恶性变。
 - ✓ NF2基因的缺乏会导致钙粘蛋白介导的细胞连接的不稳定。

・NF2基因:

- \checkmark Inactivation of NF2 has been described in many tumor types.
- Patients with neurofibromatosis type 2 syndrome are not known to develop RCCs at an increased frequency.
- Five of our 8 NF2 mutant RCC demonstrated loss of chromosome 22q where NF2 resides, consistent with biallelic inactivation of this tumor suppressor gene.

TABLE 3. NF2 Mutated Clear Cell RCC in TCGA PanCancer Atlas Database

	NF2		Tumor Size/			
Case	Variant	Age/Sex	Stage	Morphology	Outcome	Other Notable Genetic Anomalies
TCGA-CJ-4901-01	R341*	47/male	14 cm/pT3b	Clear cell	Alive 47 mo	VHL, MLH1, BAP1 deletions; loss of 3p: partial gain of 5q
TCGA-CJ-4638-01	X172_splice	46/male	15 cm/ pT3aN1M1	Clear cell	DOD 14 mo	VHL, BAP1 mutations; CDKN2A deletion; partial loss of 3p: partial gain of 5q
TCGA-B8-4153-01	D494N	74/male	5 cm/pT3a	Clear cell	Alive 25 mo	VHL, ARID1B mutations; PBRM1 deletion; partial loss of 3p
TCGA-B8-5098-01	X200_splice	53/female	4 cm/pT1	Unclassified, rhabdoid features	Died 52 mo (ESRD)	CDK12, NF1, TP53, MLH1, APC, EP400, SMARCA4 mutations: partial loss of 3p
TCGA-B0-5084-01	X121_splice	33/male	8 cm/pT3N1M1	Unclassified, possibly clear cell	DOD 7 mo	NSD1, NOTCH1 mutations
TCGA-B0-4698-01	X81_splice	75/male	10.5 cm/pT4	Unclassified with rhabdoid features	DOD 1 mo	KRAS, TP53, ARID3A, EP300 mutations

*Stop codon.

ESRD indicates end-stage renal disease.

TABLE 4. NF2 Mutated Papillary RCC in TCGA PanCancer Atlas Database

			Tumor Size/			
Case	NF2 Variant	Age/Sex	Stage	Morphology	Outcome	Other Notable Genetic Anomalies
TCGA-BQ-5875	Y132*	68/female	13 cm/pT3a	Oncocytic, reverse polarity	Alive 36 mo	SETD2, STAG2 mutations; probable 16q and 17g gain; 22g loss
TCGA-EV-5902	R359Kfs*11	58/male	3 cm/pT1N0	Biphasic, hyaline nodules	Alive 19 mo	Probable 5, 16, 20 gain
TCGA-SX-A7SL	G151Vfs*23	7/male	4 cm/pT1	Biphasic hyaline nodules	Alive 27 mo	Probable 16,17q,20 gain; 6 loss
TCGA-Y8-A896	D277Gfs*20	62/male	6 cm/pT3aNX	Unclassified but suggestive of biphasic morphology	Alive 18 mo	Probable 7 gain; 6, 9, 15q, 22q loss
TCGA-G7-6797	X200_splice	46/male	2.6 cm (multifocal)/ pT1a	Solid papillary NOS	Alive 25 mo	BAP1, MSH2, FANCD2, BCL11B mutations; probable 16, 20 gain and 1, 6p, 22q loss
TCGA-BQ-5877	K171*	60/male	3 cm/pT3aN1M1	Papillary NOS	DOD 8 mo	SETD2, BAP1 mutation; CDKN2A/B deletion; probable 12 gain and 1p,3p,11,18, 22q loss
TCGA-GL-7966	L295Rfs*14	28/female	6.5 cm/pT3N1	Tubulopapillary NOS	Alive 3 mo	<i>FH</i> and <i>B2M</i> nonsense mutations; probable partial 6p gain and 1p, 4, partial 5p, 10, 13q, 15q, 18,22q loss
TCGA-P4-A5EA X270_splic		54/female	8 cm/pT3N1	Papillary NOS	DOD 6 mo	CREBBP, MAP3K1, RASA1, BCOR nonsense or frameshift mutations
TCGA-B1-A47M	P134H	79/male	4.5 cm/pT3a	Papillary NOS	Alive 21 mo	SMARCB1, NIFE2L2, ERRFI1, TET1 mutations; probable 7 gain

*Stop codon.

DOD indicates dead of disease; NOS, not otherwise specified.

Differential Diagnosis

- ✓ Papillary RCC
- ✓ Collecting Duct Carcinoma
- ✓ MiTF Family Translocation Renal Cell Carcinoma
- ✓ Gonadal Sex Cord-Stromal Tumors
- ✓ Wilms Tumor

SUMMARY

We report a novel, morphologically distinctive subtype of RCC associated with NF2 mutations. The latter suggests the potential for targeted therapy.

THANK YOU

