

HHS Public Access

Author manuscript Genet Med. Author manuscript; available in PMC 2014 December 01.

Published in final edited form as:

Genet Med. 2013 December ; 15(12): 990-996. doi:10.1038/gim.2013.42.

Renal Growth in Isolated Methylmalonic Acidemia (MMA)

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Abstract

Purpose—We sought to predict renal growth based on clinical and metabolic parameters in patients with isolated methylmalonic acidemia (MMA), a group of disorders associated with chronic kidney disease.

Methods—Fifty MMA patients, followed from 2004 to 2011, were classified by molecular genetics and studied using a combined cross-sectional and longitudinal design that included renal ultrasound examinations, anthropometric measurements, and metabolic phenotyping. Renal length was compared to healthy controls and modeled to other clinical parameters using multiple regression analyses.

Results—Comparisons with age-matched controls showed that renal length in MMA subjects was significantly decreased (p < 0.05). Stepwise regression modeling found that combinations of height, serum cystatin C, and serum methymalonic acid concentrations best predicted kidney size. The regression equations used to generate MMA kidney nomograms were: renal length (cm) = 6.79 + 0.22 * age for the controls and 6.80 + 0.09 * age for the MMA cohort (p < 0.001; constant and slope).

Conclusions—Renal length, reflective of kidney growth, significantly decreased in MMA patients over time compared to controls and was predictable with select clinical parameters. Cystatin C and serum methylmalonic acid concentrations were highly correlated with smaller kidneys and decreased renal function in this patient population.

Keywords

isolated methylmalonic acidemia; renal growth; renal ultrasound; chronic kidney disease; cystatin C; methylmalonyl-CoA mutase; vitamin B12

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Supplementary information is available at the Genetics in Medicine website

Background

Methylmalonic acidemia (MMA) is an inborn error of metabolism caused by deficiency of the mitochondrial enzyme, methylmalonyl-CoA mutase (*MUT*). This vitamin B12 dependent enzyme catalyzes the isomerization of L-methylmalonyl-CoA into succinyl-CoA at the terminal step of propionyl-CoA oxidation.¹ Isolated MMA exhibits genetic heterogeneity and most commonly is caused by mutations in the *MUT* gene, resulting in either complete (*mut*⁰) or partial (*mut*⁻) loss of enzymatic activity, but can also be caused by *cblA* (*MMAA*), *cblB* (*MMAB*), and the much rarer *cblD variant* 2 (*MMADHC*) complementation classes. The latter forms of MMA are the result of enzymopathies that impair the intracellular transport and metabolism of cobalamin.¹ Unlike *mut* MMA, most patients with *cblA* and 50% of *cblB* forms of MMA are responsive to parenteral vitamin B12, a clinical and biochemical determinant that has been associated with improved survival and milder disease.²⁻⁷

MMA patients exhibit a well-recognized phenotype that features massive elevations of disease-related metabolites in the body fluids and multisystemic complications such as growth impairment, pancreatitis, metabolic stroke of the basal ganglia, intellectual disability and end stage kidney disease.¹ Although this disorder has historically had a poor prognosis, with most vitamin B12-nonresponsive patients dying in the first decade of life²⁻⁴, earlier diagnosis through newborn screening and more aggressive treatment has afforded increased survival.^{7,8} This highlights the need to understand the natural history of MMA and to develop improved therapies.

Chronic kidney disease is a serious complication of MMA, developing in most *mut* and *cblB* patients by the age of 8 years.^{5,6} Although more commonly associated with vitamin B12 non-responsive forms of MMA, patients with *mut*⁻ and *cblA* subtypes also can develop renal insufficiency and ultimately require kidney transplantation.^{7,9,10} While the mechanisms of renal injury in MMA have not been elucidated, the few reported kidney biopsies have consistently demonstrated tubular injury and interstitial fibrosis with secondary glomerulosclerosis.¹¹⁻¹⁴ The primary pathological manifestation of kidney injury in MMA may be mediated through mitochondrial dysfunction in the proximal renal tubule, as has been described in a *Mut* mouse model.¹⁵

The most common marker of renal function, serum creatinine, correlates poorly with glomerular function in patients with MMA. Walter et al. documented that some MMA patients had a markedly depressed glomerular filtration rate (GFR) in the setting of normal serum concentrations of creatinine.¹³ Cystatin C based estimates of GFR, which are less dependent upon muscle mass, have not yet been evaluated in the MMA population.¹⁶⁻²⁰ While the direct measurement of glomerular filtration function with agents such as inulin and iohexol is more accurate than GFR estimates, such testing is not routinely performed in MMA patients even though a small study conducted in the late 1990s documented reduced Cr-EDTA based GFR measurements in very young children with MMA. Although these authors did not describe other biochemical, imaging or enzymatic studies in the patient cohort, their observations highlight the need to develop better disease-associated biomarkers

and to assess the utility of non-invasive imaging techniques to monitor and predict kidney growth in affected children⁷.

Renal size has been proven to be a valuable and useful parameter in evaluating renal growth in normal populations and those with genetic disorders.^{21,22} Furthermore, studies in healthy controls and patients with kidney disease have repeatedly documented the correlation between renal length and other markers of renal size such as renal volume, and GFR.²³⁻²⁵ Using ultrasonographic measurements derived from a large single center cohort of patients with MMA, we examined kidney growth and assessed the contribution of selected clinical and laboratory variables to renal size using regression analyses. The predictive equations and kidney size nomograms we have developed should provide a clinical tool to help define parameters associated with disease progression in MMA, afford prognostication for long-term management and help assess renoprotective interventions in the future.

Materials and Methods

Clinical studies

Clinical and laboratory investigations were conducted through NIH study 04-HG-0127 "Clinical and Basic Investigations of Methylmalonic Acidemia and Related Disorders" (clinicaltrials.gov identifier: NCT00078078) in compliance with the Helsinki Declaration. All subjects or their guardians gave informed assent or consent. Subjects were evaluated during the period 2004 to 2011. Analysis included 84 renal ultrasound studies, each imaging the right and left kidney, from 50 non-transplanted patients aged 2.2 years to 36.3 with enzymatic and mutation confirmed forms of MMA. We treated each ultrasound from subjects with consecutive studies as a separate data point instead of averaging each patient's data because of the significant time interval, typically greater than one year, between visits. Patients were excluded after they had undergone a kidney or combined liver-kidney transplantation. The MMA patients were compared to a published control group of 209 patients from U.S. medical centers aged 0 to 19 years with indications for ultrasound that included urinary tract infections, enuresis, aniridia, scoliosis, and undescended testicle; this approach has been used previously.²⁶

Clinical and laboratory measurements

Data included height, weight, age as well as concentrations of serum creatinine, serum methylmalonic acid, cystatin C, and urine methylmalonic acid. All laboratory testing was done by standard measures through either the Department of Laboratory Medicine at NIH or Mayo Medical laboratories. Patients were subdivided into *mut, cblA*, and *cblB* groups based on cellular complementation and molecular testing. Body surface area (BSA) was calculated using the formula:²⁷ BSA(m²) = ([Height(cm) * Weight(kg)]/3600)^{1/2}. Estimated glomerular filtration rate (eGFR) for children <18 years of age was calculated using updated Schwartz equations that include serum creatinine, or serum creatinine and cystatin C, respectively. ¹⁶ For adults, the CKD-EPI Creatinine and CKD-Epi Creatinine–Cystatin C eGFR equations were used.¹⁷

Ultrasonography and laboratory measurements

The Philips iU22 xMATRIX Ultrasound system with the C5-1 1-5 MHz bandwidth transducer and the General Electric LOGIQ E9 with the C-1-5-D 1-5 MHz bandwidth transducer were used to obtain abdominal ultrasound studies for all study participants. Renal length was measured in the longitudinal axis. The original measurements were made in the context of research, and read by one of three radiologists.

Statistical analyses

Demographic variables included age, genotype, and gender. The paired *t*-test was used to compare right and left kidney length. Pearson's correlation was used to determine variation between independent variables. Published data from age-matched controls²⁶ were compared with MMA study patients using a two-sample *t*-test of the means assuming unequal variances and by a linear regression model that included the average of the right and left kidneys as the dependent variable and age as the independent variable.

In addition to variation with age, we examined the correlation of renal length with height because MMA patients can be shorter than healthy individuals. A second linear regression model was created with height as the independent variable. This linear model was compared to a different control group that created a nomogram of right and left renal length versus height using measurements from 325 German children ages 3 days to 16 years without known kidney disease.²⁸

Using renal length as the dependent variable, the following independent variables were included in a multiple regression equation: age, height, weight, body surface area (BSA), body mass index (BMI), serum methylmalonic acid (MMA_S), urine methylmalonic acid (MMA_U), serum creatinine, cystatin C, eGFR creatinine (creatinine based), eGFR creatinine - cystatin C (creatinine and cystatin C based), gender and enzymatic subtype (complementation class).

The best model, as determined by the highest correlation coefficient (\mathbb{R}^2), was developed using stepwise regression analyses. Individual predictor variables were eliminated if the regression coefficient (β) had a statistical significance *of p* > 0.05 using the Wald test. Beta weights were calculated to assess the unique contribution of independent variables relative to the regression model (beta weight is the change in renal length when an independent variable increases one standard deviation while other independent variables are held constant). Stata version 11 (StataCorp LP, College Station, TX) was used for data manipulation and analysis.

When two or more variables were closely correlated (multicollinearity), such as MMA_S and serum creatinine (Pearson coefficient of correlation 0.808 (p < 0.0001); See Table S1), they were not included in same multiple regression model.²⁹ Multicollinear relationships were observed between the variable group of creatinine, cystatin C and MMA_S, and when controlling for height, the R² values for these predictors were similar (Table S2). Height, BSA, and weight also demonstrated multicollinearity when controlling for height, with R² values of 0.767, 0.757, and 0.712, respectively.

Results

Demographics and MMA subtypes

Ultrasound measurements were available for 58 patients with isolated MMA who participated in our study. The average age of patients was 11.9 years and the age range was 2.2 years to 36.3 years. After excluding 7 patients who had kidney transplants and one patient with distorted renal anatomy due to severe scoliosis, 50 patients (86.2 %) were evaluated for further study. Of these patients, 35 were *mut* subtype yielding 63 ultrasound studies, 9 *cblA* subtype (14 ultrasound studies), and 6 *cblB* subtype (7 studies). A single kidney study includes an ultrasound of both right and left kidneys. Eighteen patients had sequential ultrasound studies yielding an additional 33 kidney studies.

Differences in renal length by anatomy, subtype, and gender

In the entire study group, the mean left kidney length was 0.37 cm greater than the right (p< 0.001); similar differences were noted in studies of healthy children.³⁰ Based on these findings, the data were divided into left and right kidneys for subsequent multiple regression analyses. The *mut* patients had slightly but not significantly smaller kidneys on average compared to the *cblA* and *cblB* subtypes, and the enzymatic subtype was insignificant when added into the regression analysis (controlling for height and cystatin C): *mut* (p = 0.487), *cblA* (p = 0.208), and *cblB* (p = 0.485). Therefore, renal data from all subtypes was pooled. Likewise, gender was not a significant variable (p = 0.554).

Renal length: correlations with age and height

Figure 2(a) and Figure 2(b) are nomograms showing renal length variation with age and height, respectively. For more severely affected patients, with serum methylmalonic acid levels greater than 2000 micromolar, renal lengths were smaller across the age and height spectrum (Figure 2).

When comparing to a control group, MMA patients had a significant difference in average renal length (p < 0.05), as shown in Table 1 and Figure 1.²⁶ The regression equation for the control group was renal length (cm) = 6.79 + 0.22 * age (years) for children older than 1 year. The linear regression equation for MMA subjects was 6.80 + 0.09 * age (years) with p < 0.001 for both the constant and β_{age} coefficient. In addition to age (R² = 0.40), renal length varied linearly with height (R² = 0.51) [Figure 2 (b)] and BSA (R² = 0.52). Because the published controls averaged right and left kidney lengths when developing their nomogram, we used an average of the right and left kidneys for comparison.²⁶

With height as the independent variable and renal length as the dependent variable, MMA patients had a diminished slope compared to historical controls, indicating that in MMA there is a decreased renal growth rate that is independent of height.²⁸ For this group of published controls, renal length nomograms were developed for left and right kidneys separately.²⁸ The control group equations were: left kidney length (cm) = 0.051 * height (cm) + 1.8, and right kidney length (cm) = 0.033 * height (cm) + 4.0, and right kidney length (cm) = 0.037 * height + 3.0 (p < 0.001 for β coefficients).

Renal length correlations with serum and urine laboratory values

Serum creatinine, MMA_S and MMA_U were analyzed according to enzymatic subtype and age (Table S3). For the *mut* and *cblA* groups, serum creatinine and MMA_S increased with age compared to MMA_U, which remained unchanged over time. The number of patients in the *cblB* group was too small to see trends similar to the *mut* and *cblA* groups. Table S1 shows that creatinine, MMA_s and MMA_U significantly (p < 0.05) correlated to age with Pearson's correlation coefficients of 0.503, 0.235, and -0.256, respectively for all enzymatic subtypes. Figure S1 shows serum creatinine, MMA_s and MMA_U variations with age. In more severely affected patients with a serum methylmalonic acid of 2000 micromolar or greater, serum creatinine rises faster with age [Figure S1(a)]; conversely, MMA_s vs. age has a greater slope when correlated with eGFR creatinine greater than 60 mL/min/1.73m² [Figure S1(b)].

Routine measurement of serum cystatin C began midway through the study and 33 of the 84 ultrasounds having matched serum cystatin C values. Cystatin C (Table S1) correlated closely with serum creatinine and MMAs and had Pearson correlation coefficients of 0.790 and 0.768, respectively. We also analyzed MMA_S compared to eGFR creatinine - cystatin C and found that MMA_S rises sharply when the eGFR creatinine - cystatin C is below 60 mL/min/ $1.73m^2$ (Figure 3).

Multiple regression modeling

The most predictive multiple regression model for renal length found cystatin C and height to be the optimal independent variables with beta weights of -0.51 and 0.75, respectively. Figure S2 is a representation of this regression model showing that renal length is inversely proportional to cystatin C when stratified by height. The final predictive equation for left kidney length (cm) = 4.9 - 0.92*cystatin C + 0.036*height; for the right kidney, predicted length (cm) = 3.6 - 0.92*cystatin C + 0.043*height. Table S2 compares MMA_S, cystatin C, creatinine, and eGFR based multiple regression models stratified by different creatinine levels. The regression model using MMA_S was found to be highly predictive of renal length when serum creatinine was below 1.5 (Table S2).

Discussion

Chronic kidney disease due to progressive tubulointerstitial injury is emerging as a frequent manifestation in MMA and eventually results in the need for dialysis or kidney transplantation: half of our cohort of MMA patients of all ages and 100% of our *mut*⁰ patients over age 15 either have been transplanted or have an eGFR below 60 mL/min/1.73 m². Kidney disease in this population is notoriously difficult to diagnose and monitor early in life, as reliance on serum creatinine-based eGFR estimates may over-state the true GFR due to low muscle mass¹³ and kidney biopsy, the most accurate test for determining the nature and extent of renal disease, is invasive and is rarely offered to MMA patients. Once clinically manifest, renal disease in MMA patients has a relentless progression to kidney failure as is illustrated by the linear decline of eGFR in the patient population, with a faster rate of decline apparent in those with more severe metabolic parameters (Figure 4). The failure to detect chronic kidney disease early is certain to delay the implementation of

renoprotective strategies in affected infants and children. Whereas blockade of the reninangiotensin system (RAS) is the current standard of care for adults with chronic kidney disease, experience with renoprotective measures in the pediatric population has not been as thoroughly studied.³ For example, there has only been one published article addressing renoprotection with RAS blockade in MMA³¹ despite the formal recognition that such studies should be pursued.³² The successful small molecule treatments of related metabolic disorders that feature tubulointerstitial disease and progressive kidney failure, such as nephropathic cystinosis and tyrosinemia type I,³² and striking preclinical efficacy of systemic AAV gene delivery to treat the kidney disease of MMA in a mouse model provides strong practical and theoretical support for the study of renoprotective interventions in MMA patients.³³

To predict renal length in MMA subjects, we first created a multivariate model and compared our data to healthy controls in the medical literature using age as the independent variable. Renal length was chosen as the dependent variable from other ultrasongraphic measurements such as renal volume and cortical thickness because renal length has the least amount of inter-observer variation and has been established to correlate with GFR.^{24,34} Compared to published healthy controls, most age groups in the MMA cohort had significantly smaller renal lengths (Table 1). The healthy control group and the MMA group both had similar constants (y-intercept), 6.79 and 6.80 respectively, suggesting that kidney size was normal at birth in the MMA patients; however, the slope (cm/year) of kidney length increase in the MMA group was less than half that of the controls: 0.09 cm/year versus 0.22 cm/year.²⁶ In comparison to another large study of 778 healthy children less than 18 years of age, the MMA kidney growth slope was only a third of the controls: 0.09 cm/year versus 0.3 cm/year.³⁵ These data provide a marker of the natural history of kidney growth in MMA, one that might be used in future studies to examine the effects of renoprotective regimens and perhaps gene therapy on the renal phenotype seen in the patients.

We were also interested in the comparison of MMA patients to healthy control subjects when height was the independent variable, especially because MMA patients have been shown to be shorter than age matched healthy subjects^{4,36}. Although most of the standardized nomograms created for renal length use age as the independent variable, one study of healthy children developed a linear model with height as the independent variable, one study of healthy children developed a linear model with height as the independent variable.²⁸ The slope of the nomogram for kidney length vs. height in the MMA patients was less than this particular control group, with right and left kidney MMA slopes of 0.037 cm (renal length)/cm (height) and 0.033 cm/cm, respectively, and control slopes of 0.050 cm/cm and 0.051 cm/cm, respectively. Thus, the comparison of MMA patients to yet another set of healthy children via linear modeling when controlling for height also demonstrated that MMA patients have slower renal growth, even when controlling for height.

Using multivariate modeling, we have found that over 70% of the variability (\mathbb{R}^2) in kidney length was captured by our model. In order to gauge how our multivariate predictive capability (\mathbb{R}^2) compared to those previously reported, we demonstrated a predictive capability of an independent multivariate model for a much larger study of 3031 healthy children under the age of 13 which found over 60% of variability was explained by their

model (R^{2} 's of 0.696 and 0.712 for right and left kidney, respectively).³⁷ Cystatin C and height were the most predictive independent variables in the multiple regression model. Table S2 shows that cystatin C, MMA_S, or serum creatinine had similar R^{2} values when controlling for height and illustrates the significance of these variables as independent predictors of renal length.

Our results establish that for serum creatinine <1.5 mg/dL, MMA_S is a stronger predictor of kidney length in the multivariate model than either cystatin C or eGFR creatinine - cystatin C (Table S2). When beta weights are included in the regression analysis, MMA_s proves to be a highly significant independent variable with a value of -0.42. The proportional relationship between MMA_S and smaller renal size in this study combined with a large beta weight for MMAS suggests that elevated MMAS is strongly associated with renal disease in methylmalonic acidemia when controlling for height by multiple regression. Additionally, Figure 2(a) and Figure 2(b) demonstrate smaller renal size versus age and height, respectively, in patients with serum methylmalonic acid greater than 2000 micromolar. The association between MMA_S and renal failure is also evident in Figure 3 where MMA_S is inversely related to eGFR creatinine - cystatin C below 60 mL/min/1.73 m²; additionally, all of our patients with a serum methylmalonic acid level above 1000 micromolar have an eGFR creatinine below 60 mL/min/1.73 m². These observations would advocate for the routine measurement of serum methylmalonic acid concentrations in the longitudinal monitoring of patients with MMA and highlight the need for future studies to examine serum markers concomitant with a measured GFR.

To complement serum creatinine, we obtained cystatin C levels with 33 of the total 84 renal ultrasounds but added this measurement after the protocol began. Using the aggregate data, the multivariate model incorporating cystatin C emerged as the most precise model, explaining over 76% of renal length variation ($R^2 = 0.767$) when controlling for height (see Figure S2). As expected, the regression model using cystatin C was more predictive than the model using GFR estimates calculated from serum creatinine and cystatin C (Table S2), which is most likely due to the muscle mass differences in MMA patients and the patients used to derive the eGFR equation.^{16,17} Our data suggests that serum cystatin C concentration should also be included in the routine monitoring of renal function in MMA patients and warrants further study, including validation against direct GFR assessments, especially because recent studies suggest that in the general population, GFR estimates that incorporate both serum creatinine and serum cystatin C appear to be the most accurate.^{18,19} In a population with reduced muscle mass such as MMA, the GFR equation using cystatin C or cystatin C and MMAs may be preferred.

Prior investigations have not consistently demonstrated a relationship between MMA_U and kidney disease.^{5,13} A multicenter, retrospective review of 35 patients suggested that MMA_U predicted chronic kidney disease as determined from calculations using the Schwartz formula; however, inter- and intra-individual variations were claimed to contribute to overlap in the enzymatic subgroups of this study.⁵ In contrast, Walter et al. found no correlation between random urinary methylmalonic acid levels and reduction in GFR.¹³ We found that the urine methylmalonic acid concentration was not a significant predictor of renal length when controlling for height (p = 0.471): MMA_U did not correlate with trends in

serum creatinine as can be seen by the statistically insignificant Pearson's correlation in Table S1 (R = 0.0389). Given the poor correlation between MMA_U, and renal length or trends in serum creatinine, our data advocates using MMA_S as a predictor of renal disease and including this analyte in the routine monitoring of patients with MMA.

The *mut* patients had slightly smaller kidneys than the other subgroups; however, when controlling for cystatin C and height, we did not detect a significant difference between enzymatic subtypes on kidney length. Another study found an increase in chronic kidney disease in *mut*⁰ and *cblB* compared to other enzymatic subtypes causing isolated MMA, which would suggest that these individuals might have smaller kidneys.⁵ A possible explanation for this discrepancy between the studies may derive from the small number of participants with *cblB*; in both our study (n=6) and that of Horster et al. (n=9).⁵ In addition to MMA subtypes, we did not find gender to be a contributor to renal length in MMA patients. Boys' and girls' body sizes grow at different rates; however, multiple studies, including ours, have demonstrated that renal length is not significantly different between the sexes.³⁰

Our study has a number of limitations. Despite using the largest single center MMA patient cohort described to date, the sample size was comparatively small and we were unable to use a discovery and replication design. Second, the majority of our data was cross-sectional, which may introduce increased variability in the nomograms compared to data derived from a purely longitudinal design. Third, the most severely affected patients may not have been fit for travel to NIH, and thus underestimated our effect size.

In summary, the rate of renal growth in the MMA subjects studied here was one half to one third that observed in controls.^{26,35} Renal length is likewise significantly decreased compared to normal controls and predicted by a multiple regression model that uses the readily available clinical variables of height and cystatin C. When serum creatinine values fall in the age normal range, MMA_S and cystatin C are comparable predictors of renal length, suggesting that both should be included in the routine evaluation in MMA patients. In addition to generating a clinically useful nomogram to follow kidney growth in MMA, our results capture a marker of the natural history of the renal disease of MMA and should assist with the study of therapeutic interventions designed to target kidney disease in this population.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

This work was supported by the Intramural Research Programs of the NHGRI and NIDDK, NIH.

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Figure 1.

Renal length (average of left and right) vs. age in methylmalonic acidemia (MMA) and agematched controls²⁶. Mean \pm SD. Asterisk denotes p < 0.05.

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Figure 2.

(a). Renal length vs. age, stratified by serum methylmalonic acid level. Patients with greater than 2000 micromolar serum methylmalonic acid have smaller kidneys. (b). Renal length vs. height, stratified by serum methylmalonic level. Similar to Figure 2(a), more severely affected patients with higher serum methylmalonic acid levels have smaller kidneys across height spectrum.

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Figure 3.

Serum methylmalonic acid (MMAS) for all enzymatic subtypes vs. eGFR creatinine cystatin C. Best fitted linear regression lines are shown for eGFR = 60 and eGFR > 60. MMA_S increases markedly for eGFR values below 60 mL/min/1.73 m².

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Figure 4.

eGFR creatinine for all enzymatic subtype patients declines with age, reflective of a progressive chronic kidney disease. eGFR calculated using the updated Schwartz equation for children.¹⁶ For adults, the CKD-EPI creatinine equation was used.¹⁷ The solid line represents linear regression for patients with serum methylmalonic acid less than 2000 micromolar and dashed lines represent a linear regression for serum methylmalonic acid greater than 2000. The patients with a higher serum methylmalonic acid show a lower eGFR.

Renal Length of MMA subjects vs. Controls, by age

Table 1

The average and SD for renal length values for MMA patients (all enzymatic subtypes) and control subjects are presented in one year age intervals. Right and left renal lengths are averaged. Comparisons using the t test assumed unequal variances. Data from controls are from Rosenbaum et al.²⁶ Asterisk denotes data not available.

	MMA Patie	ints		Controls	
Age (years)	Observations	Renal Length (cm)	Observations	Renal Length (cm)	<i>p</i> -value
2 to 3	9	6.22 ± 0.88	12	7.36 ± 0.54	0.01
3 to 4	9	7.19 ± 0.78	30	7.36 ± 0.64	0.32
4 to 5	7	6.88 ± 0.80	26	7.87 ± 0.50	0.013
5 to 6	2	6.48 ± 1.17	30	8.09 ± 0.54	0.00
6 to 7	7	7.24 ± 1.2	14	7.83 ± 0.72	0.13
7 to 8	9	7.87 ± 0.86	18	8.33 ± 0.51	0.13
8 to 9	4	8.39 ± 0.99	18	8.90 ± 0.88	0.19
9 to 10	2	7.5 ± 1.77	14	9.20 ± 0.90	0.19
10 to 11	9	7.57 ± 1.17	28	9.17 ± 0.82	0.008
11 to 12	3	7.64 ± 0.67	22	9.60 ± 0.64	0.008
12 to 13	2	9.03 ± 0.53	18	10.42 ± 0.87	0.023
13 to 14	5	9.12 ± 0.62	14	9.79 ± 0.75	0.039
14 to 15	2	8.85 ± 0.07	14	10.05 ± 0.62	<0.001
15 to 16	*	*	9	10.93 ± 0.76	*
16 to 17	1	8.65	10	10.04 ± 0.86	*
17 to 18	3	9.52 ± 0.42	4	10.53 ± 0.29	0.016
18 to 19	1	8.8	8	10.81 ± 1.13	*
TOTAL	62		768		